STIC Search Report Biotech-Chem Library



STIC Database Tracking Number: 119952

From: Paul Schulwitz

Location: Biotech-Chem Library

REM-1A65

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TO: James Schultz Location: REM-2D18/2C18

Art Unit: 1635

Wednesday, April 21, 2004 Case Serial Number: 10/001844

Examiner Schultz,

Search Notes

See attached results.

If you have any questions about this search feel free to contact me at any time.

Thank you for using STIC search services!

Paul Schulwitz Technical Information Specialist STIC Biotech/Chem Library (571)272-2527



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AAV18416 ADB00919 PCR primer Shh-D s RT-PCR primer Shh-PCR primer for mou Rat Shh coding seq Human MDZ3 scannin

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Human Acetylcholin Human ALDHS allele Human/mouse C/EBP Human Sonic hedgeh						ID NO:43.	Sonic hedgehog; Shh; polymer conjugate; polyalkene glycol group; higavailihiltv: formulation: neurological disorder:	, 7,1	neurodegenerative disorder; Parkinson's disease; Huntington's disease;	Alzheimer's disease; neurological injury; stroke; multiple sclerosis;	1 tumour;	
	ST.					imer, SEQ	polyalken	der, cance	disease;	y; stroke;	oectoderma	
AAS96144 ABA99313 ABA02229 AAF27039	ALIGNMENTS		BP.			Human Sonic hedgehog (Shh) mutagenic primer, SEQ ID NO:43	r conjugate;	inflammatory disorder; autoimmune disorder; cancer;	Parkinson's	ogical injur	malignant glioma; medulloblastoma; neuroectodermal tumour;	
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Garber 99US-0137011P. 99US-0149016P. Pepinsky RB, Taylor F, WPI; 2001-049927/06. (BIOJ ) BIOGEN INC. 01-JUN-1999; 13-AUG-1999;

26-MAY-2000; 2000WO-US014741.

WO200073337-A1. Homo sapiens. Synthetic.

07-DEC-2000.

Modified hedgehog protein, useful in the treatment of Parkinson's disease and Huntington's chorea, comprises a polymer containing a polyalkylene glycol group linked to any residue other than the N-terminal and lysine residues. 

Example 6; Page 77; 157pp; English.

The invention relates to novel polymer conjugates of hedgehog proteins which have increased bloavailability. The hedgehog proteins are conjugated to a non-naturally-occurring polymer complyinging a polyalkylene glycol group, with the proviso that the polymer is not conjugated to the protein used in the conjugate may be a wild-type or mutant sonic hedgehog protein used in the conjugate may be a wild-type or mutant sonic hedgehog protein used in the conjugate may be a wild-type or mutant sonic hedgehog (Shh), Indian hedgehog (Ihh) or Desert hedgehog (Dhh) protein. The hedgehog (Ihh) or Desert hedgehog (Dhh) protein, or may be a hedgehog fusion protein. The invention also relates to methods of a federal mapping functionally important regions of a protein by modifying accessible amino acid side chains, and determining the effect the position and/or type of modification have on the activity of the protein. The hedgehog polymer conjugates may be used in the management of protein. The hedgehog polymer conjugates may be used in the management of uniquence and autoimmum diseases, and cancers. In particular, they may be used to prevent preventing or ameliorate neurodegenerative of disorders (e.g., Parkinson's disease, Huntington's disease, Alzheimer's disease); age-associated neurodogical disease, neurological injury and trauma; immunological diseases of the nervous system (e.g., multiple selent in increased half-life, altered tissue distribution (such as may result in increased half-life, altered tissue distribution

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an improved ability to stay in the vasculature for longer periods of time), increased stability in solution, protection from protectory degradation, or reduced immunogenicity. In particular, the ability to remain in the vasculature for prolonged periods may allow a hedgehog protein of the invention to cross the blood-brain barrier, and an increased thermal stability would be an advantage when formulating the hedgehog protein in powder form. The present sequence represents a human sould be an exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                invention
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Sequence 38 BP; 8 A; 11 C; 9 G; 10 T; 0 U; 0 Other;

Gaps ö Ouery Match

8.5%; Score 36.4; DB 1; Length 38;
Best Local Similarity 97.4%; Pred. No. 0.031;
Matches 37; Conservative 0; Mismatches 1; Indels 162 GACTGGGTGTACTACGAGTCCAAGGCACATATCCACTG 199 ð

38 GACTGGGTGTACTACGAGTGCAAGGCACATATCCACTG 1

025/c AAF27025 standard; DNA; 49 BF. AAF27025; RESULT 

(first entry) 30-MAR-2001 Human Sonic hedgehog (Shh) mutagenic primer, SEQ ID NO:29.

Sonic hedgehog; Shh; polymer conjugate; polyalkene glycol group; bioavailibility; formulation; neurological disorder; infilammatory disorder; autoimmune disorder; cancer; neurodegenerative disorder; Parkinson's disease; Huntington's disease; Alzheimer's disease, neurological injury; siroke; multiple sclerosis; malignant glioma; medulloblastoma; neuroectodermal tumour;

Homo sapiens Synthetic. WO200073337-A1.

07-DEC-2000

26-MAY-2000; 2000WO-US014741.

01-JUN-1999; 99US-0137011P. 13-AUG-1999;

(BIOJ ) BIOGEN INC.

Pepinsky RB, Taylor F,

Garber E;

WPI; 2001-049927/06.

Modified hedgehog protein, useful in the treatment of Parkinson's disease and Huntington's chorea, comprises a polymer containing a polyalkylene glycol group linked to any residue other than the N-terminal and lysine residues.

Example 2; Page 67; 157pp; English.

The invention relates to novel polymer conjugates of hedgehog proteins which have increased bicavailability. The hedgehog proteins are conjugated to a non-naturally-occurring polymer comprising a polyalkylene glycol group, with the proviso that the polymer is not conjugated to the N-terminus, or to lysine residues of the hedgehog protein. The hedgehog protein used in the conjugate may be a wild-type or mutant Sonic hedgehog (Shh), Indian hedgehog (Shh) or Desert hedgehog (Dhh) protein, or may be a hedgehog fusion protein. The invention also relates to methods of defining and mapping functionally important regions of a protein by

modifying accessible amino acid side chains, and determining the effect
the position and/or type of modification have on the activity of the
protein. The hedgehog polymer conjugates may be used in the management of
various medical conditions including various neurological disorders,
inflammatory and autoimmune diseases, and cancers. In particular, they
any be used to prevent preventing or ameliorate neurodegenerative
cd isorders (e.g., Parkinson's disease, Huntington's disease, Alzheimer's
disease); age-associated neurological disease, neurological injury and
traums; immunological diseases of the nervous system (e.g., multiple
cd disease); seroke; and malignant gliomas, medulloblastomas and
cc neuroectodermal tumours. The modifications made to the hedgehog protein
cc neuroectodermal tumours. The modifications made to the hedgehog protein
cc nimproved ability to stay in the vasculature for longer periods of
c time), increased stability in solution, protection from proteolytic
cdegradation, or reduced immunogenicity. In particular, the ability to
cermain in the vasculature for prolonged periods may allow a hedgehog
cc protein of the invention to cross the blood-brain barrier, and an
cincreased thermal stability would be an advantage when formulating the
chedgehog protein in powder form. The present sequence represents a human
cc increased mutagenic primer used in an exemplification of the

Sequence 49 BP; 8 A; 18 C; 9 G; 14 T; 0 U; 0 Other;

ö 8.5%; Score 36; DB 1; Length 49; 88.6%; Pred. No. 0.067; tive 0; Mismatches 5; Indels o.57 Best Local Similarity 88.67 Matches 39, Conservative

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AAF27038 standard; DNA; 39 AAF27038/c RESULT 3

30-MAR-2001 (first entry)

AAF27038;

Human Sonic hedgehog (Shh) mutagenic primer, SEQ ID NO:42.

Sonic hedgehog; Shh; polymer conjugate; polyalkene glycol group; bloavallibility; formulation; neurological disorder; inflammatory disorder; autoimmune disorder; cancer; neurodegenerative disorder; Parkinson's disease; Huntington's disease; Alzheimer's disease; neurological injury; stroke; multiple sclerosis; malignant glioma; medulloblastoma; neuroectodermal tumour; mutagenic primer; ss. 

Homo sapiens. Synthetic.

WO200073337-A1.

07-DEC-2000.

26-MAY-2000; 2000WO-US014741.

99US-0137011P. 99US-0149016P. 01-JUN-1999; 13-AUG-1999;

BIOJ ) BIOGEN INC.

Garber E; Pepinsky RB, Taylor F,

WPI; 2001-049927/06.

Modified hedgehog protein, useful in the treatment of Parkinson's disease and Huntington's chorea, comprises a polymer containing a polyalkylene glycol group linked to any residue other than the N-terminal and lysine residues.

Example 6; Page 77; 157pp; English.

The invention relates to novel polymer conjugates of hedgehog proteins which have increased bloavailability. The hedgehog proteins are conjugated to a non-naturally-occurring polymer comprising a polyalkylene conjugated to a non-naturally-occurring polymer comprising a polyalkylene glycol group, with the proviso that the polymer is not conjugated to the N-terminus, or to lysine residues of the hedgehog protein. The hedgehog protein. The hedgehog protein in the conjugate may be a wild-type or mutant sonic hedgehog (Shh), indian hedgehog (Inh) or besert hedgehog protein. The mapping functionally important regions of a protein by modifiving accessible amino acid side chains, and determining the effect the position and/or type of modification have on the activity of the position and/or type of modification have on the activity of the protein. The hedgehog polymer conjugates may be used in the management of various medical conditions including various neurological disorders, inflammatory and autoimmune diseases, and cancers. In particular, they may be used to prevent preventing or ameliorate neurodegenerative disorders (e.g., Parkinson's disease, Huntington's disease, Alzheimer's disease, immunological diseases, muchington's disease, Alzheimer's disease, immunological diseases, medulioblastomas and traumay immunological diseases, medulioblastomas and cancercine, increased sasociated neurological disease, medulioblastomas and carcodermal tumours. The modifications made to the hedgehog protein may result in increased half-life, altered tissue distribution (such as an improved ability to stay in the vasculature for longer periods of time), increased stability in solution, protection from protectory can improved ability to stay in the vasculature for prolonged periods may allow a hedgehog creased the invention to cross the bload-brain barrier, and an increased thermal stability would be an advantage when ferents a human company in the degehog mutagenic primer used in an exemplification of the

Sequence 39 BP; 7 A; 12 C; 13 G; 7 T; 0 U; 0 Other;

Gaps ö Score 35.8; DB 1; Length 39; Pred. No. 0.043; 2; Indels 97 CCACGTCTGACCGCGACCGCAGCAAGTACGGCATGCTGG 135 н 39 CCACGICTGACCGCGATCGCTGCAAGTACGGCATGCTGG 0; Mismatches 8.4%; Local Similarity 94.9 les 37, Conservative Query Match Matches ઠ 셤

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Sonic hedgehog; Shh; polymer conjugate; polyalkene glycol group; bicavailbbility; formulation; neurological disorder; inflammatory disorder; autoimmune disorder; cancer; neurodegenerative disorder; parkinson's disease; huntington's disease; halzheimer's disease; neurological injury; stroke; multiple sclerosis; malignant glioma; medulloblastoma; neuroectodermal tumour; Human Sonic hedgehog (Shh) mutagenic primer, SEQ ID NO:41. AAF27037 standard; DNA; 37 BP (first entry) mutagenic primer; ss 30-MAR-2001 AAF27037; AAF27037/c RESULT 

WO200073337-A1. Homo sapiens. Synthetic.

26-MAY-2000; 2000WO-US014741. 07-DEC-2000

99US-0137011P. 1999;

useful in the treatment of Parkinson's disease Garber 99US-0149016P protein, F, Taylor WPI; 2001-049927/06. (BIOJ ) BIOGEN INC. Modified hedgehog 13-AUG-1999; Pepinsky RB,

The invention relates to novel polymer conjugates of hedgehog proteins which have increased bioavailability. The hedgehog proteins are which have increased bioavailability. The hedgehog proteins are conjugated to a non-naturally-occurring polymer comprising a polyalkylene glycol group, with the proviso that the polymer is not conjugated to the protein used in the conjugate may be a wild-type or mutant Sonic hedgehog (Shh), Indian hedgehog (Inh) or Desert hedgehog (Dhh) protein. The hedgehog (Inh) or Desert hedgehog (Dhh) protein, or may be a hedgehog qualon protein. The invention also relates to methods of defining and mapping functionally important regions of a protein by modifyling accessible amino acid side chains, and determining the effect the position and/or type of modification have on the activity of the protein. The hedgehog polymer conjugates may be used in the management of various medical conditions including various neurological disoaces, and cancers. In particular, they may be used to prevent preventing or ameliorate neurodegenerative disorders, or protein memorogeniated neurological disease, huntington's disease, Alzheimer's disease); stroke; and malignant gliomas, medicallobical injury and cancers in memoroccodermal tumours. The modifications made to the hedgehog protein may result in increased half-life, altered tissue distribution (such as mimproved ability to stay in the vasculature for longer periods of time), increased stability in solution, protection from proteolytic degradation, or reduced immunogenicity. In particular, the ability to remain in the vasculature for prolonged pariods and an expensive may and an expensive may exceed the proper prolonged pariods may allow a hedgehog protein the process of the prolonged pariods may allow a hedgehog protein may result in the vasculature for prolonged pariods may allow a hedgehog process. protein of the invention to cross the blood-brain barrier, and an increased thermal stability would be an advantage when formulating the hedgehog protein in powder form. The present sequence represents a human sonic hedgehog mutagenic primer used in an exemplification of the and Huntington's chorea, comprises a polymer containing a polyalkylene glycol group linked to any residue other than the N-terminal and lysine Example 6; Page 77; 157pp; English. 

Sequence 37 BP; 6 A; 10 C; 12 G; 9 T; 0 U; 0 Other;

Gaps ö 7.9%; Score 33.8; DB 1; Length 37; 44.6%; Pred. No. 0.099; ve 0; Mismatches 2; Indels 94.68; 35; Conservative Similarity Query Match Best Local S

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38 CGAAGATGGCCACCACTCAGAGGAGTCTCTGCACTAC 74 CGAAGATGGCCACCACTGCGAGGAGTCTCTGCACTAC 37 ઠે 셤

AAF27041/c ID AAF27041 standard; DNA; 35

RESULT 5

AAF27041;

Sonic hedgehog; Shh; polymer conjugate; polyalkene glycol group; bioavailibility; formulation; neurological disorder; inflammatory disorder; aucoimmune disorder; cancer; neurodegenerative disorder; Parkinson's disease; Huntington's disease; Alzheimer's disease; neurological injury; stroke; multiple sclerosis; malignant glioma; medulloblastoma; neuroectodermal tumour; Human Sonic hedgehog (Shh) mutagenic primer, SEQ ID NO:45. (first entry) 30-MAR-2001 

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Modified hedgehog protein, useful in the treatment of Parkinson's disease and Huntington's chorea, comprises a polymer containing a polyalkylene glycol group linked to any residue other than the N-terminal and lysine
                                                                                                                                              Example 6; Page 77; 157pp; English.
                                                  26-MAY-2000; 2000WO-US014741
                                                                                          Pepinsky RB, Taylor F,
mutagenic primer; ss
                                                                                                      WPI; 2001-049927/06
                                                                               (BIOJ ) BIOGEN INC.
                            WO200073337-A1
            Homo sapiens.
                                                             01-JUN-1999;
                                       07-DEC-2000.
                  Synthetic
                                                                                                                                    residues
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The invention relates to novel polymer conjugates of hedgehog proteins which have increased bioavailability. The hedgehog proteins are which have increased bioavailability. The hedgehog proteins are conjugated to a non-naturally-occurring polymer comprising a polyarlkylene glycol group, with the proviso that the polymer is not conjugated to the protein used in the conjugate may be a wild-type or mutant Sonic hedgehog protein used in the conjugate may be a wild-type or mutant Sonic hedgehog (Shh), Indian hedgehog (Ihh) or Desert hedgehog (Dhh) protein or may be a hedgehog fusion protein. The invention also relates to methods of a hedgehog fusion protein. The invention also relates to methods of the conjugate may be used in the management of the position and/or type of modification have on the activity of the protein. The hedgehog polymer conjugates may be used in the management of various medical conditions including various neurological disorders, inflammatory and autoimmune diseases, and cancers. In particular, they may be used to prevent preventing or amellorate neurodegenerative disorders, of inflammatory and autoimmune diseases, and cancers. In particular, and malignant glionas, meduloblastoms and contrological injury and trauma; immunological diseases of the nervous system (e.g., multiple contenum, immunological diseases of the nervous system (e.g., multiple may result in increased half-life, altered tissue distribution (such as maniguation, or reduced immunogenicity. In particular, the ability to carmain in the vasculature for longer periods of the invention or reduced immunogenicity. In particular, harrier and an expense or the invention or reduced immunogenicity. In particular, and an entition or remain in the vasculature for prolonged periods may allow a hedgehog process of the invention of the inventi process of the invention to cross the blood-brain barrier, and an increased thermal stability would be an advantage when formulating the hedgehog protein in powder form. The present sequence represents a human sonic hedgehog mutagenic primer used in an exemplification of the invention

Sequence 35 BP; 8 A; 15 C; 9 G; 3 T; 0 U; 0 Other;

Gaps ö ch 7.8%; Score 33.4; DB 1; Length 35; l Similarity 97.1%; Pred. No. 0.1; 34; Conservative 0; Mismatches 1; Indels Query Match Best Local Si Matches 34;

139 GCCTGGCGGTGGAGGCCGGCTTCGACTGGGTGTAC 173 35 GCCTGGCGGTGGAGGCCTGCTTCGACTGGGTGTAC

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AAF27040 standard; DNA; 37 BP AAF27040/c ID AAF2704 XX RESULT 6

(first entry) 30-MAR-2001 AAF27040; 

Human Sonic hedgehog (Shh) mutagenic primer, SEQ ID NO:44.

Sonic hedgehog; Shh; polymer conjugate; polyalkene glycol group; bloavailibility; formulation; neurological disorder; inflammatory disorder; autoimmune disorder; cancer; neurodegenerative disorder; parkinson's disease; Muntington's disease, Alzheimer's disease; neurological injury; stroke; multiple sclerosis; malignant glioma; medulloblastoma; neuroectodermal tumour; mutagenic primer; ss.

Homo sapiens Synthetic. WO200073337-A1.

Garber E;

99US-0137011P

07-DEC-2000.

26-MAY-2000; 2000WO-US014741.

99US-0137011P. 99US-0149016P. 01-JUN-1999; 13-AUG-1999;

(BIOJ ) BIOGEN INC.

ä Garber Pepinsky RB, Taylor F,

WPI; 2001-049927/06.

Modified hedgehog protein, useful in the treatment of Parkinson's disease and Huntington's chorea, comprises a polymer containing a polyalkylene glycol group linked to any residue other than the N-terminal and lysine

Example 6; Page 77; 157pp; English.

The invention relates to novel polymer conjugates of hedgehog proteins which have increased bioavailability. The hedgehog proteins are which have increased bioavailability. The hedgehog proteins are conjugated to a non-naturally-occurring polymer comprising a polyalkylene glycol group, with the proviso that the polymer is not conjugated to the proviso that the polymer is not conjugated to the proviso that the polymer is not conjugated to the protein used in the conjugate may be a wild-type or mutant sonic hedgehog protein. The hedgehog [Ihh] or Desert hedgehog protein. The hedgehog [Ihh] or Desert hedgehog protein, or may be cefining and mapping functionally important regions of a protein by condition and/or type of modification have on the activity of the position and/or type of modification have on the activity of the protein. The hedgehog polymer conjugates may be used in the management of various medical conditions including various neurological disease, and cancers. In particular, they cardinate the prevent preventing or amelicate neurodegenerative disease), age-associated neurological disease, Huntington's disease, Alzheimer's disease, jaca-associated neurological disease, incurological linjury and cracers in munological tiseases of the nervous system (e.g., multiple callerose), stroke, and malignant gliomas, medulloblastomas and cancercodermal tumours. The modifications made to the hedgehog protein may result in increased half-life, altered tissue distribution (e.g. multiple called in moreased stability in solution, protection from periods of time), increased stability in solution, protection from periods may allow a hedgehog cental in the vasculature for prolonged periods may allow a hedgehog contends in the heavell the normal advantance the normal and and and and allogablastine and and and and and allogablastine and and and and and allogablastine and and and and and and allogablas a human be an advantage when formulating the increased thermal stability would be an advantage when formulating hedgehog protein in powder form. The present sequence represents sonic hedgehog mutagenic primer used in an exemplification of the nvention

Sequence 37 BP; 7 A; 8 C; 13 G; 9 T; 0 U; 0 Other;

Query Match

7.6%; Score 32.2; DB 1; Length 37;

RESULT

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The sequences given in AAQ91654-57 are primers which were used to amplify a sequence which encodes a human sonic hedgehog protein, homologous to a brosophila hedgehog protein (AAR77337). The human sequence was isolated by screening of human genome DNA by nested polymerase chain reaction using these primers, followed by use of a clone to screen a human fetal lung 5'-stretch plus cDNA library in phage lambda-gtl0. A clone has been isolated from a phage Pl library in phage lambda-gtl0. A clone has been primers SHHF (AAQ91654) and SHHF (AAQ91655), to give clone SHHP1. A 2.5-kb ECORI CA repeat fragment is amplified using primers SHHCAR (AAQ91666) and SHHCAR (AAQ91667). Probes and primers derived from the sonic hedgehog sequence may be used as diagnostic agents for neuromuscular, autonomic or central nervous system disorders, and the gene may also be used in gene therapy. A puripodies generated from the encoded protein may be used as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sonic hedgehog; SHH gene; HH gene; tumorigeneais; oncogeneais; basal cell carcinoma; breast cancer; medulloblastoma; tumour; cell proliferation; cell differentiation; diagnosis; therapy; human; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hedgehog-like protein(s) and nucleic acid(s) encoding them - useful to treat degenerative nervous system disorder(s) and in gene therapy.
             Human, sonic hedgehog gene; nested polymerase chain reaction; PCR; fetal lung; probe; primer; diagnostic; nervous system disorder; gene therapy; antibody; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human mutated sonic hedgehog (SHH) gene exon 2 PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5.6%; Score 24; DB 1; Length 24;
100.0%; Pred. No. 3.8;
cive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 24 BP; 6 A; 5 C; 11 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                               (HARD ) HARVARD COLLEGE.
(IMCR ) IMPERIAL CANCER RES TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24 ACCGAGGGCTGGGACGAGGATGGC 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 ACCAAGGCTGGGACGAAGATGGC 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 5; Page 100; 210pp; English.
                                                                                                                                                                                                                                                                                                                                                                                             Ingham PW, Mcmahon AP, Tabin CJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         therapeutic or research reagents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP
                                                                                                                                                                                                                            94WO-US014992.
                                                                                                                                                                                                                                                                    93US-00176427
94US-00356060
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAV18405 standard; cDNA; 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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Matches 24; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1995-255060/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-SEP-1998
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                                                                                                                                             WO9518856-A1
                                                                                                                                                                                                                            30-DEC-1994;
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14-DEC-1994;
                                                                                                                                                                                    13-JUL-1995.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        primer; ss
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                                                                                                     Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to a method for determining the presence of neoplastic molecular markers in a host, involving the use of neoplastic molecular marker specific reagents to detect such markers and analysing the array of reagents, allowing the identification of the neoplastic disease present. This can be used to determine the best treatment for cancers, in paritudiar neural cell, lung and prostate tumours. The present sequence is a PCR primer useful for detecting the coding sequences of markers of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Determining the presence of neoplastic molecular markers, by identifying the presence of markers in host test sample using array of neoplastic molecular marker specific reagents and analyzing the array of the
                                                                                                                                                                                                                                                                                                                                                                       Human; cancer; neoplastic disease; tumour specific marker; cytostatic; transcription factor; PCR; primer; ss.
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                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6.3%; Score 27; DB 1; Length 27; 100.0%; Pred. No. 1.2; 1.1ve 0; Mismatches 0; Indels
                        3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 27 BP; 3 A; 11 C; 9 G; 4 T; 0 U; 0 Other;
                                                           38 CGAAGATGGCCACCACTCAGAGGAGTCTCTGCACTAC 74
                                                                                          CGAAGATGGCCACCACTCATGCGAGTCTCTGCACTAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human sonic hedgehog protein gene primer SHHFS'.
  91.9%; Pred. No. 0.21; tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     255 TCGGCCACGGTGCACCTGGAGCAGGC 281
                                                                                                                                                                                                                                                                                                                                    Human SHH gene PCR primer SEQ ID NO: 289
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 19; 41pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAQ91654 standard; cDNA; 24
                                                                                                                                                                                                          ABT03768 standard; DNA; 27
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  Best Local Similarity 91.9
Matches 34; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CEMI-) CEMINES ITC
                                                                                                                                                                                                                                                                                          13-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
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                                                                                                                                                                                                                                                    ABT03768;
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Gaps

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AAQ91654 RESULT

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This human sonic hedgehog (SHH) gene exon 2-specific primer was used with another exon 2-specific primer (see AAV18406) in a PCR using DNA from Muman bacterial artificial chromosome (BAC) DNA pools. Only pools comprising a BAC that contains the sequence tag defined by the primer pair will yield an amplification product. The process was continued until as ingle positive BAC was identified. The positive clone, BAC270A17, was dispessed with restriction enzymes and ligated into vectorette linkers. Mutations (see AAV18403 and AAV18404) have been identified in the SHH gene in human cancers. The mutated SHH genes and the encoded polypeptides (see AAW48735 and AAW48736) can be used in methods for the treatment and diagnosis of cancer and other diseases involving cell proliferation or
                                                                                                                                                                                                                                                    New nucleic acid encoding oncogenic human hedgehog protein - useful for, e.g. treatment and diagnosis of cancer and diseases involving cell proliferation or differentiation.
                                                                                                                                                                                                                                                                                                                                    Example; Page 23; 47pp; English.
                                                                                                                                                                          Hu Z, Bonifas J;
                                                        97WO-US020227
                                                                                            96US-00748591
                                                                                                                                    (REGC ) UNIV CALIFORNIA
                                                                                                                                                                                                                WPI; 1998-297857/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  differentiation
                                                        12-NCV-1997;
                                                                                              13-NOV-1996;
                     22-MAY-1998.
                                                                                                                                                                             Epstein E,
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5.6%; Score 24; DB 1; Length 24; 100.0%; Pred. No. 3.8; tive 0; Mismatches 0; Indels Sequence 24 BP; 6 A; 5 C; 11 G; 2 T; 0 U; 0 Other; 24 ACCGAGGGCTGGGACGAGGATGGC 47 Acceaegerregaceaegargec 24 24; Conservative Local Similarity Query Match g à

Gaps

AAD10171 standard; DNA; 24 BP (first entry) 12-SEP-2001 AAD10171; RESULT 10 AAD1017.

Human, Sonic hedgehog, Shh; morphogenic signal; neuron; embryonic patterning; cell culture; cell differentiation; ischaemia; cell proliferative disorder; intracerebral grafting; Huntington's chorea; neurological disorder; Alahaimar's disease; Parkinson's disease; amyotrophic lateral sclerosis; ALS; multiple sclerosis; PCR primer; ss. Human Sonic hedgehog (Shh) gene amplifying forward PCR primer SHHF 93US-00176427. 94US-00356060. 95US-00435093. 95US-00460900. 96US-00674509 04-MAY-1995; 05-JUN-1995; 05-JUN-1995; sapiens JS6261786-B1 02-JUL-1996; 17-JUL-2001, Ношо 

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The present invention relates to assay for screening compounds that corporate or inhibit binding of hedgehog polypeptide to naturally concentring patched receptor. The hedgehog proteins comprise morphogenic signals produced by embryonic patcarning centres, and are involved in the formation and maintenance of ordered spatial arrangements of differentiated tissues in vertebrates, both adult and embryonic. The proteins can be used to generate and/or maintain a array of different vertebrate tissues both in vitro and in vivo. The invention also relates to ammanian cell (e.g. neuron, testicular cell) responsive to hedgehog induction. Hedgehog agonists and anicanance of neurons and various vertebrate organogenic pathways. The hedgehog gene is useful in cell culture techniques to enhance survival and maintenance of neurons and various vertebrate organogenic pathways. The hedgehog gene is useful in determining whether a patient is at the risk of disorder characterised by unwanted cell proliferation or aberrant control of differentiation. The ceptage proteins can be used to induce foctal neurons especially neuronal stem cells in intracerebral grafting. The protein or its mimetic can be used in the treatment of neurological conditions e.g. cinjury to nervous system, isochaemia resulting from stroke, Alzheimer's condisease, Parkinson's disease, Hunthington's chorea, amyotrophic lateral conditions 
                                                                                                                                                                                                                   Screening compounds that potentiate or inhibit binding of hedgehog polypeptide to naturally occurring patched receptor, comprises contacting polypeptide with receptor and test compound, and detecting change in
                                                                                                 Mcmahon AP
(IMCR ) IMPERIAL CANCER RES TECHNOLOGY LTD. (HARD ) HARVARD COLLEGE.
                                                                                                                                                                                                                                                                                                                                                                                                 Example 5; Col 98; 127pp; English
                                                                                                 Ingham PW,
                                                                                                 Tabin CJ,
                                                                                                                                                               WPI; 2001-440859/47
                                                                                                 Marigo V,
                                                                                                                                                                                                                                                                                                   polypept
binding.
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0; Gaps 5.6%; Score 24; DB 1; Length 24; 100.0%; Pred. No. 3.8; ive 0; Mismatches 0; Indels 47 24 ACCGAGGGTGGGACGAAGATGGC Local Similarity 100. Query Match Best Loc Matches 8

Sequence 24 BP; 6 A; 5 C; 11 G; 2 T; 0 U; 0 Other;

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Hedgehog protein, sonic hedgehog; Shh; indian hedgehog; Ihh; desert hedgehog; cell differentiation; human; PCR primer; 88. Human Shh DNA amplifying primer SHHF5'. 1 Acceaecereceaeceaeceaece 24 AAH76132 standard; DNA; 24 29-OCT-2001 (first entry) Homo sapiens. US6271363-B1 AAH76132; RESULT 11 AAH76132 g

93US-00176427. 94US-00356060. 95US-00435093. 97US-00954698

30-DEC-1993; 14-DEC-1994; 04-MAY-1995;

20-OCT-1997; 07-AUG-2001.

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                                                                                                                                                             The invention relates to nucleic acids encoding hedgehog proteins selected from sonic hedgehog (Shh), indian hedgehog (Ihh), desert hedgehog (Dh) polypeptides. The hedgehog genes are involved in the formation of ordered spatial arrangements of differentiated tissue in vertebrates. The nucleic acid sequences are useful for producing hedgehog proteins, used for promoting differentiation of, or survival of differentiation on unronal calls, and for promoting proliferation, survival or differentiation of mesenchymal, endodermal or ectodermal tissue, particularly chondrocytes, or testicular germ line cells. Sequences AAH76132-133 represent PCR primers for amplifying a human Shh DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hedgehog related-protein, sonic hedgehog protein, Shh, ischemia, stroke, desert hedgehog protein, Dhh, indian hedgehog protein; Ihh, neuron, neurological condition, nervous system injury, tumour-induced injury, aging, Alzheimer, s disease; chronic neurodegenerative disease; Parkinson's disease; thronic neurodegenerative disease; spinocerebellar degeneration, chronic immunological disease;
                                                                                              Novel nucleic acid encoding a hedgehog polypeptide, used to produce the polypeptide, which is used to promote proliferation, survival, and/or differentiation of neuronal and mesodermal tissue.
                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bumcrot DA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primer for cDNA encoding human sonic hedgehog protein (Shh).
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                                                                                                                                                                                                                                                                                                            Score 24; DB 1; Length 24;
Pred. No. 3.8;
                                                                                                                                                                                                                                                                                                                                  0; Indels
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                                                                                                                                                                                                                                                                                       Sequence 24 BP; 6 A; 5 C; 11 G; 2 T; 0 U; 0 Other;
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100.0%; Pred. No. ...
0; Mismatches
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(IMCR ) IMPERIAL CANCER RES TECHNOLOGY LTD.
                              IMCR ) IMPERIAL CANCER RES TECHNOLOGY LID.
                                                                                                                                                                                                                                                                                                                                                       24 ACCGAGGCTGGGACGAAGATGGC 47
                                                                                                                                                                                                                                                                                                                                                                           1 Acceaegecreeaceaegargec 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          multiple sclerosis; PCR primer; ss
                                                    Tabin CJ;
                                                                                                                                          Example 5; Col 88; 118pp; English.
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94US-00356060.
95US-00435093.
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95US-00462386
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-APR-2001 (first entry)
                                                                                                                                                                                                                                                                                                                      Local Similarity 100.
                                                      Mcmahon AP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ingham PW, Mcmahon AP,
                     (HARD ) HARVARD COLLEGE
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                                                                          WPI; 2001-456723/49
05-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
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04-MAY-1995;
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                                                      Ingham PW,
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                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 12
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PCR primers AAC87097-98 were used to amplify cDNA encoding a hedgehog related-protein. The specification describes a sonic hedgehog protein (ShN), a desernt hedgehog protein (DhN), and an indian hedgehog protein (IN)). The hedgehog polynucleotides are useful in diagnostic, in antisense therapy and in therapeutic assays for detecting and treating disorders involving, e.g., aberrant expression of vertebrate hedgehog compared in the polynetides are useful therapeutically to enhance conditions deriving from acute, subacute, or chronic injury to the conditions deriving from acute, subacute, or chronic injury, to the conditions system, including traumatic injury, chemical injury, vasal injury and deficits (such as the ischemia resulting from stroke), together with infectious/inflammatory and induced injury, aging of the nervous system including Alzheimer's disease, chronic neurodegenerative diseases of the nervous system, including Parkinson's disease, Huntington's chorea, amylotrophic lateral sclerosis, spinocerebellar degenerations, and chronic immunological diseases of the nervous system or affecting the coloresis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sonic hedgehog; Shh; desert hedgehog; Dhh; Indian hedgehog; Ihh; antiparkinsonian; antiarrhythmic; neuroprotective; anticonvulsant; extostatic; noctropic; spermatogenesis; peripheral nervous system; central nervous system; Alzheimer's disease; Parkinson's disease; Huntington's disease; arrhythmia; nerve degeneration; multiple sclerosis; immunological disorder; neoplastic; hyperplastic; PCR primer; ss.
   for treating diseases
Polynucleotides encoding hedgehog proteins, useful for treating diseases of nervous system such as Alzheimer's disease, Parkinson's disease, Huntington's chorea, amylotrophic lateral sclerosis, multiple sclerosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5.6%; Score 24; DB 1; Length 24; 00.0%; Pred. No. 3.8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 24 BP; 6 A; 5 C; 11 G; 2 T; 0 U; 0 Other;
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(IMCR ) IMPERIAL CANCER RES TECHNOLOGY LTD.
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                                                                                                                                      Example 5; Col 86; 119pp; English.
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94US-00356060.
95US-00435093.
95US-00462386.
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nes 24; Conservative
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04-MAY-1995;
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New vertebrate hedgehog-related proteins, useful e.g. for promoting differentiation, survival and proliferation of cells, e.g. for treating neurodegeneration.

Example 5; Col 88; 116pp; English.

The present invention describes an isolated and/or recombinant comprising a hedgehog (hh) amino acid (aa) sequence concoded by a nucleic acid (il) that hybridizes under stringent conditions to lof 6 sequences (see ABN87544, and ABN87546 to ABN87550). (I) binds to a natural patched receptor. Specifically claimed example of (I) are cyprentially and ABN87545 to ABN87550). (I) binds to a natural patched receptor. Specifically claimed example of (I) are cyprentially concorpic, neuroprotective, anticonvulsant, antiarrhythmic and cytostatic activities. (I) induces the expression of the BMP-2 and 4 genes, and of the How gene. (I) and be used: (i) to promite differentiation of courconal cells and survival of the Gifferentiation of comminergic or motor neurons, proliferation of chondrocytes, and compaminergic or motor neurons, proliferation of chondrocytes, and concorpinate of alteration and/or survival of meadcemal or cetcodermal cells, either in cell cultures (particularly for preparation of transplants) or therapeutically; (ii) for detecting loss of response, in tissues or, to hh proteins; (iii) in drug screening (to identify (aut) agonists, useful e.g. for inhibition of spermatogenesis), and (iv) corticat e.g. injuries/defects in the central or peripheral nervous system, e.g. multiple sclerosis, neoplastic and hyperplastic con arrhythmias caused by nerve degeneration; immunological disorders of the nervations in the central nervous system, also to promote attachment of prostheses. The present sequence represents a PCR primer for human sonic confirmation in the central nervous system, also to promote attachment of invention in the exemplification of the present

Sequence 24 BP; 6 A; 5 C; 11 G; 2 T; 0 U; 0 Other;

0; Gaps Query Match 5.6%; Score 24; DB 1; Length 24; Best Local Similarity 100.0%; Pred. No. 3.8; Matches 24; Conservative 0; Mismatches 0; Indels

24 ACCGAGGCTGGGACGAAGATGGC 47 g

RESULT 14 ADA26284

20-NOV-2003 (first entry)

Human Sonic hedgehog (Shh) cDNA PCR primer #1.

Human; PCR; ss; Sonic hedgehog; Shh; neuronal cell; skeletogenesis; chrodrogenesis; osteogenesis; degenerative disorder; nervous system; neuronal cell death; neural cell; neuromuscular disorder; nervous autonomic disorder; central nervous system disorder; anoxia; ischaemia; peripheral nervous system disorder; achycarda; atrial cardiac arrhythmia; striated heart; stem cell development; digestive tract; liver; multiple sclerosis; primer.

JS2003054437-A1.

ADA26284 standard; DNA; 24 BP. ADA26284; BEST SERVICE S

Homo sapiens.

97US-00954771. 20-OCT-1997;

93US-00176427. 94US-00356060. 95US-00435093. 30-DEC-1993; 14-DEC-1994; 04-MAY-1995;

95US-00462386 05-JUN-1995;

(INGH/) INGHAM P W. (MCMA/) MCMAHON A P. (TABI/) TABIN C J.

Tabin CJ; Ingham PW, Mcmahon AP,

WPI; 2003-555377/52

Modulating growth, differentiation or survival of a cell, useful for treating a degenerative disorder of the nervous system characterized by neuronal cell death, comprises contacting the cell with a hedgehog polypeptide.

Example 5; Page 48; 121pp; English.

The invention relates to a method for modulating growth, differentiation or survival of a cell, comprising contacting the cell with a hedgehog or survival of a cell, comprising contacting the cell to differentiate to a neuronal cell phenotype comprising contacting the cell to differentiate to a neuronal cell phenotype comprising contacting the cell with a hedgehog polypeptide, modulating skeletogenesis by contacting a target tissue and treating a degenerative disorder of the nervous system characterised by neuronal cell death, comprising calministering a hedgehog polypeptide causing prolonged survival of neural cells in the patient, relative to the absence of hedgehog treatment. The hedgehog polypeptides are useful for treating a degenerative disorder of cells in the patient, relative to the absence of hedgehog treatment. The hedgehog polypeptides are useful for treating a degenerative disorder of neuromuscular, autonomic or central nervous system disorders of neuromiscular, autonomic or central nervous system disorders of neuromal degeneration associated with a natural aging process. The specifically Alzheimer's disease, Muntington's disease, motorophic selerosis, neuronal damage resulting from anoxia, ischaemia or trauma and neuronal degeneration associated with a natural aging process. The collisorders including disorders affecting peripheral nervous system disorders including disorders affecting peripheral nervous system disorders including disorders affecting neuronal cardiac arrhythmias which may arise from a degenerative condition whereby the nerves innervate the striated muscle of the heart, in nerve prostheses for repairing central and peripheral nerve damage, for treating neoplastic or hyperplassic corpusions and in controlling the development of stem cells created promations and in controlling the development of stem cells created promation of the disease of teme cells created promation of the disease of teme cells created by the development of stem cells created by the perver and peripheral nerves s

Sequence 24 BP; 6 A; 5 C; 11 G; 2 T; 0 U; 0 Other;

.; 0 Match 5.6%; Score 24; DB 1; Length 24; Local Similarity 100.0%; Pred. No. 3.8; les 24; Conservative 0; Mismatches 0; Indels Query Match Best Local Matches

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Gaps

g

ADD25290 standard; DNA; 24 BP RESULT 15 ADD25290

15-JAN-2004 (first entry) *************

ADD25290;

Human Sonic hedgehog PCR primer #1.

hedgehog; patched receptor; spermatogenesis inhibition; ovary function inhibition; embryogenesis; differential tissue maintenance; 88; PCR; primer; human.

Homo sapiens.

US6576237-B1

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                                                                                                                                                                                                                                                                                              The invention relates to an isolated antibody (I) which is immunoreactive with a hedgehog polypeptide (II) that binds to a patched receptor, where (II) is encoded by uncleic acid which hybridise to a fully defined vertebrate hedgehog (hh) protein. (I) is useful as a hedgehog antagonist by blocking action of naturally occurring hedgehog protein, and therefore for inhibiting spermatogenesis. (I) is also useful for inhibiting normal ovarian function. (I) is useful for blocking the action of one or more hedgehog proteins and allows the study of the role of these proteins e.g., embryogenesis and/or maintenance of differential tissue. (I) is also useful in immunohistochemical staining of tissue samples in order to evaluate the abundance and pattern of expression of the hedgehog protein abundance and pattern of expression of the hedgehog protein to detect and evaluate hedgehog protein levels as a part of clinical testing procedure. The present sequence represents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human, cell differentiation, Desert hedgehog; Dth; Sonic hedgehog; shh; Indian hedgehog; Ith; skeletogenesis; degenerative disorder; ischaemia; Alzheimer's disease; Parkinson's disease; parkinson's disease; ampliple sclerosis; Pick's disease; and sight sclerosis; Pick's disease; aging process; trauma; anoxia; antisense gene therapy; neuroprotective; anticonvulsant;
                                                                                                                                                                                                                           for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                           Novel isolated antibody which is immunoreactive with a vertebrate hedgehog protein sequence that binds with patched receptor, useful blocking action of naturally occurring hedgehog protein, and for
                                                                                                                                                              Marti-Gorostiza
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 24; DB 1; Length 24;
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                                                                                                                                                              Tabin CJ, Bumcrot DA,
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(IMCR ) IMPERIAL CANCER RES TECHNOLOGY LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 100.0%; Pred. No. 3.8 les 24; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                          Example 5; SEQ ID NO 43; 120pp; English.
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                                                           93US-00176427.
94US-00356060.
95US-00435093.
95US-00460900.
                                     16-AUG-2000; 2000US-00639695.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5.68;
                                                                                                                                                                                                                                                  inhibiting spermatogenesis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAD62117 standard; DNA; 24
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                                                                                                                                                                                      WPI; 2003-799823/75.
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                                                           30-DEC-1993;
14-DEC-1994;
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           10-JUN-2003
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The present invention relates to a novel method for modulating growth, differentiation or survival of a cell. The method involves contacting the cell with a hedgehog polypeptide such as Desert hedgehog (Dhh), Sonic hedgehog (shh). The method is used to his sonic cell to differentiate to a neuronal cell phenotype. It is used to induce a modulate skeletogenesis. The method is used to treat a degenerative disorders of the nervous system such as neuronuscular, autonomic or central nervous system such as neuronuscular, autonomic or disease, amyotrophic lateral sclerosis, Huntington's disease, Parkinson's disease, amyotrophic lateral sclerosis, Wintington's disease, multiple sclerosis, Wintington's disease, and neuronal damage resulting from trauma and neuronal damage resulting from soxia-ischemata. The invention is also used for antisense gene therapy. The present sequence is human Shh DNA amplifying PCR primer. This sequence is used in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hedgehog polypeptide; tissue array generation; tissue array maintainance; hedgehog; human; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                           Modulating cell growth, differentiation or survival, for treating
neurodegenerative diseases, such as Alzheimer's or Parkinson's disease,
comprises contacting the cell with a hedgehog polypeptide.
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                                                                                                                                                                                                                                                                                               Tabin CJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 5; Page 49; Opp; English.
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                                                     93US-00176427.
94US-00356060.
95US-00435093.
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94US-00356060.
95US-00435093.
95US-00460900.
95US-00462386
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADD71413 standard; DNA; 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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                                                                                                                                                                                                                                                                                            Ingham PW, Mcmahon AP,
                                                                                                                                                                       (INGH/) INGHAM P W. (MCMA/). MCMAHON A P. (TABI/) TABIN C J.
                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-803151/75
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                                                  30-DEC-1993;
14-DEC-1994;
04-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-DEC-1993;
14-DEC-1994;
04-MAY-1995;
05-JUN-1995;
35-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-JAN-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADD71413;
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(HARD ) HARVARD COLLEGE.
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Marti-Gorostiza Bumcrot DA, Tabin CJ, Mcmahon AP, Ingham PW,

WPI; 2003-831623/77

New nucleic acid encoding a hedgehog polypeptide having an amino acid sequence identical or homologous to a vertebrate hedgehog protein, useful for generating or maintaining an array of different vertebrate tissue in vitro and in vivo.

Example 5; SEQ ID NO 43; 118pp; English.

The invention describes an isolated nucleic acid encoding a hedgehog polypeptide having an amino acid sequence identical or homologous to a vertebrate hedgehog protein or its portion and not identical to a fully defined 471-bp sequence. The nucleic acid is useful for generating and/or maintaining an array of different vertebrate tissue both in vitro and in vivo. This sequence represents a primer used to isolate DNA encoding vivo. This sequence mundant sonic hedgehog.

Sequence 24 BP; 6 A; 5 C; 11 G; 2 T; 0 U; 0 Other;

Gaps ö 5.6%; Score 24; DB 1; Length 24; 100.0%; Pred. No. 3.8; tive 0; Mismatches 0; Indels 24; Conservative Query Match Best Local Similarity Matches

## 24 ACCGAGGGCTGGGACGAAGATGGC 47

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AAV18406 standard; cDNA; 25 RESULT

AAV18406;

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(first entry) 14-SEP-1998 Human mutated sonic hedgehog (SHH) gene exon 2 PCR primer.

Sonic hedgehog; SHH gene; HH gene; tumorigenesis; oncogenesis; basal cell carcinoma; breast cancer; medulloblastoma; tumour; cell proliferation; cell differentiation; diagnosis; therapy; human; PCR; primer; ss 

Synthetic

Homo sapiens,

WO9821227-A1

12-NOV-1997;

22-MAY-1998

96US-00748591. 13-NOV-1996;

(REGC ) UNIV CALIFORNIA.

WPI; 1998-297857/26.

Bonifas J;

Hu 2,

Epstein E,

New nucleic acid encoding oncogenic human hedgehog protein - usefu e.g. treatment and diagnosis of cancer and diseases involving cell proliferation or differentiation.

Example; Page 23; 47pp; English.

used with This human sonic hedgehog (SHH) gene exon 2-specific primer was used vanother exon 2-specific primer (see AAV18406) in a PCR using DNA from

human bacterial artificial chromosome (BAC) DNA pools. Only pools comprising a BAC that contains the sequence tag defined by the primer pair will yield an amplification product. The process was continued until a single positive BAC was identified. The positive clone, BAC270A17, was digested with restriction enzymes and ligated into vectorette linkers. Mutations (see AAV18403 and AAV18404) have been identified in the SHH (see AAM48735 and AAV18404) can be used in methods for the treatment and diagnosts of cancer and other diseases involving cell proliferation or differentiation 888888888888888

Sequence 25 BP; 4 A; 8 C; 8 G; 5 T; 0 U; 0 Other;

Gaps ô Score 23.4; DB 1; Length 25; Pred. No. 5.5; 0; Mismatches 1; Indels Query Match
Best Local Similarity 96.0%;
Matches 24; Conservative

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116 CAGCAAGTACGGCATGCTGGCCCGC 140 cadcaadracedcarecreec 25

à d RESULT 19

BP. ABZ79785 standard; DNA; 24 ABZ79785 ID ABZ7

ABZ79785;

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(first entry) 12-MAY-2003 Indian hedgehog PCR primer SEQ ID NO:5.

Osteopathic; antirheumatic; antiarthritic; cytostatic; cartilage; cartilage differentiation; joint disease; bone fracture; myeloma; osteoporosis; rheumatoid arthritis; human; Indian hedgehog; PCR primer;

sapiens Omor

Synthetic. 

WO2003000870-A1.

03-JAN-2003

25-JUN-2002; 2002WO-JP006351

26-JUN-2001; 2001JP-00193503

(TAKE ) TAKEDA CHEM IND

Hikichi Y, Inazuka M;

WPI; 2003-201422/19

Culture method for cartilage differentiation from cells under hypoxic conditions into cartilage cells applicable in cartilage transplantation, and studying genes or proteins relating to joint diseases.

Sxample 3; Page 29; 37pp; Japanese.

The present invention describes a method for cartilage differentiation by culturing cells capable of differentiating into cartilage under hypoxic conditions. Also described: (1) a method for producing cartilage cells or cartilage by culturing the required cells under hypoxic conditions; (2) drugs containing the produced cartilage cells or cartilage. (3) a method for preventing or treating joint diseases by transplanting an effective amount of the cartilage cells or cartilage; (4) the use of the cartilage cells or cartilage; (4) the use of the cartilage cells or cartilage by using the call or cartilage differentiation or joint diseases; (5) a method for screening genes relating to cartilage (6) a method for screening promoters or inhibitors of cartilage differentiation by using promoters or inhibitors of cartilage differentiation by using any of the culture methods; (7) a method for screening preventive methods; (7) a method for screening preventive for joint diseases by using the culture

Sequence 19 BP; 3 A; 5 C; 6 G; 5 T; 0 U; 0 Other;

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methods; (8) drugs containing the screened promoters or inhibitors of cartilage differentiation, or preventives or remedies for joint diseases; (9) a method for preventing or treating joint diseases by administering an effective dose of the promoters or inhibitors, or preventives or remedies to mammals; and (10) the use of the promoters or inhibitors, or preventives or remedies for producing drugs for joint diseases. The produced cultured cartilage cells or cartilage can be used in cartilage transplantation, studying genes or proteins relating to joint diseases and screening drugs for their treatment, including diseases of bone fracture, myeloma, osteoporosis and rheumatoid arthritis. The present sequence represents a PCR primer for Indian hedgehog, which is used in an example from the present invention

Sequence 24 BP; 3 A; 4 C; 10 G; 7 T; 0 U; 0 Other;

Gaps .. DB 1; Length 24; 1; Indels 5.0%; Score 21.4; E 95.7%; Pred. No. 13; ative 0; Mismatches 22; Conservative Query Match Best Local Similarity Matches

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410/c AAV18410 standard; cDNA; 19 BP.

AAV18410;

(first entry) 14-SEP-1998 Human mutated sonic hedgehog (SHH) gene exon 2 PCR primer.

Sonic hedgehog; SHH gene; HH gene; tumorigenesis; oncogenesis; basal cell carcinoma; breast cancer; medulloblastoma; tumour; cell proliferation; cell differentiation; diagnosis; therapy; human; PCR; primer; ss

Homo sapiens. Synthetic

WO9821227-A1

22-MAY-1998

97WO-US020227. 12-NOV-1997; 96US-00748591. 13-NOV-1996;

(REGC ) UNIV CALIFORNIA.

Bonifas J; Epstein E, Hu Z,

WPI; 1998-297857/26.

New nucleic acid encoding oncogenic human hedgehog protein - useful for, e.g. treatment and diagnosis of cancer and diseases involving cell proliferation or differentiation. Example, Page 23; 47pp; English.

This human sonic hedgehog (SHH) gene exon 2-specific primer was used with another exon 2-specific primer (see AAV18410) in a PCR amplification of genomic DNA from 34 independent basal cell carcinomas. 18 4 medulloblastomas and 6 breast carcinomas. PCR primers (see AAV18407-08 and AAV18411-12) specific for SHH exons 1 and 3 were also used. PCR and AAV1841s. 2 Mutations (see AAV18403 and AAV18404) were identified in the SHH gene from 4 human cancers. The mutated SHH genes and the encoded by Porppetides (see AAV48735 and AAV1875) can be used in methods for the treatment and diagnosis of cancer and other diseases involving cell proliferation or differentiation AAV19410/ 110 AAV10/ AAV10/

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                                                                                                                                                           Sonic hedgehog; SHH gene; HH gene; tumorigenesis; oncogenesis; basal cell carcinoma; breast cancer; medulloblastcma; tumour; cell differentiation; diagnosis; therapy; human; PCR;
                 Gaps
                 .
Query Match 4.5%; Score 19; DB 1; Length 19; Best Local Similarity 100.0%; Pred. No. 23; Matches 19; Conservative 0; Mismatches 0; Indels
                                                                                                                                            Human mutated sonic hedgehog (SHH) gene PCR primer.
                                 194 CCACTGCTCGGTGAAAGCA 212
                                            CCACTGCTCGGTGAAAGCA 1
                                                                                           AAV18416 standard; cDNA; 19 BP
                                                                                                                                                                                                                                                                                                                Epstein E, Hu Z, Bonifas J;
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                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                               (REGC ) UNIV CALIFORNIA.
                                                                                                                                                                                                                                                                                                                              WPI; 1998-297857/26.
                                                                                                                                                                                                              Homo sapiens.
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                                                                                                                           14-SEP-1998
                                                                                                                                                                                                                             WO9821227-A1
                                                                                                                                                                                                                                              22-MAY-1998
                                                                                                                                                                                      primer; ss.
                                                                                                                                                                                                      Synthetic
                                                                                                           AAV18416;
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ö cDNA derived from human epidermal keratinocytes was amplified by 3-stage nesting using sonic hedgehog (5HH) gene stage 1 primers (see AAV18413 and AAV18414).

AAV184114), stage 2 primers (see AAV18415 and AAV18416) and stage 3 primers (see AAV18415 and AAV18415). The PCR product was identified as authentic SHH, A single somatic mutation (see AAV18403) of the SHH gene was found in cancers arising from 3 different tissues in independent patients. Another mutation (see AAV18404) was identified in another cancer. The mutated SHH genes and the encoded polypoptides (see AAV48735) and AAV48736) can be used in methods for the treatment and diagnosis of cancer and other diseases involving cell proliferation or differentiation Gaps .. 0 4.5%; Score 19; DB 1; Length 19; [00.0%; Pred. No. 23; 0; Indels Sequence 19 BP; 2 A; 6 C; 5 G; 6 T; 0 U; 0 Other; 100.0%; Pred. No. 284 CACCAAGCTGGTGAAGGAC 302 19; Conservative Similarity Query Match Local Best Loca Matches δ

New nucleic acid encoding oncogenic human hedgehog protein - useful for, e.g. treatment and diagnosis of cancer and diseases involving cell

e.g. treatment and diagnosis of c proliferation or differentiation.

Example; Page 25; 47pp; English.

22 RESULT

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The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ12, MDZ12. MDZ3 is encoded at chromosome 7422.1. MDZ4 is encoded at chromosome 6921.3-22.2. MDZ7 is encoded at chromosome 6921.3-22.2. MDZ4. MDZ4. MDZ4. MDZ2, encoded at chromosome 6923. MDZ4, MDZ4, MDZ4, MDZ3, MDZ4, MDZ1, or MDZ12. The mucleic caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The mucleic caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The mucleic alcids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are proteins are useful as therapeutic agents for gene therapy or as protein are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
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                                                                                                                                                                                 Cytostatic; immunostimulant; gene therapy; vaccine; human;
zinc finget protein; MD23; MD24; MD27; MD212; chromosome 7q22.1;
chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer;
developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New zinc finger-containing proteins and nucleic acids, useful in
manufacturing a medicament for treating or preventing a disorder
associated with decreased or increased expression or activity of MDZ3,
MDZ4, MDZ1 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                            Human MDZ3 scanning oligonucleotide SEQ ID 1905,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         th 4.4%; Score 18.6; D
Similarity 84.0%; Pred. No. 53;
21; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 8; SEQ ID NO 1905; 103pp; English.
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                     BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Shannon M, Gu Y, Nguyen C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    02-AUG-2001; 2001US-00922181
                     ADB00919 standard; DNA; 25
                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (AEOM-) AEOMICA INC
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                                                                                                                                                                                                                                                                                      Homo sapiens.
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                                                                                                   20-NOV-2003
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                                                              ADB00919;
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Best Local
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ACI66417
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ADB00919
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The invention discloses a microarray comprising a plurality of nucleic acid probes including one of 2,018,500 fully defined sequences, or its conscious including one of 2,018,500 fully defined sequences, or its conscious disclosed is a method of gene expression analysis. The array is used in monitoring gene expression levels by hybridisation to a DNA library, in analysis of genetic variation or in hybridisation to a DNA library, or manlysis of genetic acid probes are specifically designed for analysis of at least one target sequence. The method of analysis comprises of at least one target sequence. The method of analysis comprises or more nucleic acid probes and detecting the hybridisation. The nucleic acid probes are attached to a solid support. The analysis comprises monitoring probes are attached to a solid support. The analysis comprises monitoring gene expression levels, identifying biallelic markers or polymorphisms, or family membors of a gene and a cross-species comparison. Each of the nucleic acids further comprises a tag sequence. The array of nucleic acids further comprises a tag sequence. The array of mucleic acids further comprises a tag sequence. The array of mucleic acid further comprises a tag sequence. The array of mucleic acid further comprises a tag sequence or specific contactions of any gene, in mapping the 5' termini of mRNA molecules by primer extensions or in screening cDNA or genomic libraries or subclones of for additional subclones containing segments of DNA that have been contained and previously sequenced in the microarray. Note: The sequence of ata for this patent can also be obtained in electronic format directly from them instructions of the sequence of the format directly at each are sentenced.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           New array of nucleic acid probes, useful for in situ hybridization, in Southern, Northern or dot-blot hybridization to identify or detect the sequence or specific mutations of any gene.
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                EST; ss; probe; expressed sequence tag; microarray; gene expression; genetic variation; biallelic marker; polymorphism; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 25 BP; 7 A; 8 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   rom USPTO at segdata.uspto.goc/sequence.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human Desert hedgehog gene sense PCR primer.
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4.0%; Pred. No. 53;
.ve 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; SEQ ID NO 66408; 9pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 CGACGACAAGTAGGTCTTCGAC 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           386 CGACGCCCAAGAAGGTCTTCTAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAV62410 standard; DNA; 20 BP
                                                                                                                                                                                                                                            15-MAR-2002; 2002US-00098263.
                                                                                                                                                                                                                                                                                       16-MAR-2001; 2001US-0276759P.
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nes 21; Conservative
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                                                                  cross-species comparison
                                                                                                                                                                                                                                                                                                                               (AFFY-) AFFYMETRIX INC
                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-567953/53.
                                                                                                                                                     US2003104410-A1.
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                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                            Mittmann MP;
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Matches
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2X2X5X8X
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Human microarray DNA oligonucleotide SEQ ID NO 66408.

(first entry)

14-OCT-2003

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This sense primer corresponds to nucleotides 460-479 of a cDNA clone (see AAV62396) coding for novel human Desert hedgehog protein (see AAW79596).

I was used with an antisense primer (see AAV62411) in a first-steep PCR amplification of human leukaemia plasma cell line ARH-77 (ATCC CRL-1621) cDNA in a modified PCR method of 3'RACE. 2 Subsequent PCR amplifications (see AAV62423-26) yielded a cDNA clone (see AAV6239) encoding a C-terminal fragment (see AAW79599) of the novel human Desert hedgehog protein. Nucleotide sequences (see AAV62339-5) encoding mature and The Desert hedgehog DNA, protein and a claimed monoclonal antibody can be used in to elucidate hereditary morphological abnormalities in humans to establish their treatments and diagnoses
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCR primer, neuroectoderm cell; cell production; Parkinson's disease; early primitive ectoderm-like cell; EPL cell; cell therapy; transgenic animal; gene therapy; neuronal disease; Huntington's disease; lysosomal storage disease; multiple sclerosis; memory disorder; behavioural disorder; Alzheimer's disease; organ transplant;
                                                                                                                                                                                                                                                                                                Human Desert hedgehog protein - and corresponding DNA and monoclonal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4.3%; Score 18.4; DB 1; Length 20; 95.0%; Pred. No. 35; 1; Indels cive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 3 A; 5 C; 6 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                   (HAYB ) HAYASHIBARA SEIBUTSU KAGAKU.
                                                                                                                                                                                                                                                                                                                                           Example 1-4; Page 10; 39pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         156 GGCTTCGACTGGGTGTACTA 175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 GGCTTCGACTGGGTCTACTA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          behavioural disorder; Alzheim: spinal cord disorder; Shh; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAF87046 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12-JAN-2001; 2001WO-AU000030.
                                                                                                                         98EP-00303187
                                                                                                                                                      97JP-00121578.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR primer for Shh gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                 WPI; 1998-544642/47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     VO200151611-A1.
                             Homo sapiens.
                                                                                                                                                      25-APR-1997;
14-APR-1998;
                                                                                                                         24-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-JUL-2001
                                                          EP874048-A2.
                                                                                       28-0CT-1998
                                                                                                                                                                                                                                   Ariyasu T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAF87046;
              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                 antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 25
AAF87046/
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This sequence represents a PCR primer for the Shh gene, used within the scope of the invention. The invention relates to a method for producing scope of the invention. The invention relates to a method for producing concrected early primitive ectoderm—like (EPL) cells and a neural-inducing conditioned primitive ectoderm—like (EPL) cells and a neural-inducing conditioned confident to a mine sufficient to generate controlled confident to a mine sufficient to generate controlled confident to a nimal cells or partially differentiated progeny are differentiation to (I). The cells or partially differentiated progeny and pendual gene thereapy, the screening of pharmaceutical that induce a biological response in neuroectoderm cells or their partially differentiated progeny and evaluation of feature cells. The method is useful for producing controlled for meural cells. The method is useful for producing controlled for maintaining neuroectoderm cells. The method can cell populations. It can also be used for producing content of neuroal diseases, including Parkinson's disease, content cells and be used for producing content of neuroal diseases, including Parkinson's disease.

Contention's disease, lysosomal storage diseases, multiple sclerosis, also be used for produced by the method are used for the treatment of spinal correction of cells produced by the method are used for the treatment of spinal corrections.
                                                                                                                                                                             Producing neuroectoderm cells for treatment of Parkinson's and Alzheimer's and for transplantation comprises culturing early primitive ectoderm-like cells in conditioned medium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4.3%; Score 18.4; DB 1; Length 20; 95.0%; Pred. No. 35; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ds; Hedgehog protein; cancer; PCR; primer; amplification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Seguence 20 BP; 5 A; 2 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hedgehog protein derivative primer 2.
                                                                                                                                                                                                                                                        Example 3; Page 41; 91pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    355 ACAGCGACTICCICACTITC 374
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14-JAN-2000; 2000AU-00005098.
20-APR-2000; 2000AU-00007045.
27-APR-2000; 2000AU-00007143.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           95.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                        Rathjen J;
                                                                       (BRES-) BRESAGEN LTD
                                                                                                                                           WPI; 2001-432908/46.
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Matches 19; Conserv
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                                                                                                        Rathjen PD,
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Gaps

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Wed Apr

The primers AAV59457-V59462 were used in the production of hedgehog (hh) protein derivative may be used in the prediction and diagnosis various diseases e.g. cancer Hedgehog protein derivative and gene encoding it and diagnosis of various diseases e.g. lung cancer. Sequence 25 BP; 3 A; 9 C; 8 G; 5 T; 0 U; 0 Other; Disclosure; Page 6; 7pp; Japanese ဗ (ASAG ) ASAHI GLASS WPI; 1998-499061/43. 

Score 18.2; DE
Pred. No. 64;
0; Mismatches 244 24 gradedeceaaareeaaceere 2 222 GIGGCGGCCAAATCGGGAGGCTG Match 4.3%; Local-Similarity 87.0%; les 20; Conservative ( Query Match Best Local-Si Matches 20, à

ADB00921 standard; DNA; 25 BP Human MDZ3 ADB00921; RESULT 27 ADB00921 

scanning oligonucleotide SEQ ID 1907. (first entry) 20-NOV-2003

Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.

Homo sapiens

EP1281758-A2

05-FEB-2003.

30-JUL-2002; 2002EP-00016874

02-AUG-2001; 2001US-00922181.

(AEOM-) AEOMICA INC

Nguyen Gu ₹, Shannon M,

ΰ

WPI; 2003-423107/40.

MDZ3, New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ4, MDZ7 or MDZ12, e.g. cancer.

Example 8; SEQ ID NO 1907; 103pp; English.

The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MD23, MD24, MD27, MD212. MD23 is encoded at chromosome 7g22.1, MD24 is encoded at chromosome 6p21.3-22.2, MD27 is encoded at chromosome 15g26.1. The MD23, MD24, MD21, and MD212 is encoded at chromosome 15g26.1. The MD23, MD24, MD21, and MD212 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MD23, MD24, MD27, or MD212, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for disgnosing or monitoring a disease caused by altered expression of MD23, MD24, MD27, or MD212. The nucleic

probes are acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes ar useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention. 888888888

Sequence 25 BP; 3 A; 11 C; 4 G; 7 T; 0 U; 0 Other;

useful for prediction

Gaps ö Query Match
4.3%; Score 18.2; DB 1; Length 25;
Best Local Similarity 87.0%; Pred. No. 64;
Matches 20; Conservative 0; Mismatches 3; Indels

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363 ITCCTCACTITCCTGGACGGCGA 385 rrccrcacrarccraccccccca 23 н

> ò g

a Of

RESULT 28 ADB00920

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Gaps

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Length 25; 3; Indels

DB 1;

BP. 25 ADB00920 standard; DNA;

ADB00920;

(first entry) 20-NOV-2003

Human MDZ3 scanning oligonucleotide SEQ ID 1906.

Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MD23; MD24; MD27; MD212; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.

Homo sapiens

EP1281758-A2

05-FEB-2003.

30-JUL-2002; 2002EP-00016874.

02-AUG-2001; 2001US-00922181

(AEOM-) AEOMICA INC

Gu Y, Nguyen C; Shannon M,

WPI; 2003-423107/40.

MDZ3 New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ4, MDZ7 or MDZ12, e.g. cancer.

Example 8; SEQ ID NO 1906; 103pp; English.

The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ1, MDZ4, MDZ7, MDZ12. MDZ3 is proteins and their coding sequences: MDZ3, MDZ4, MDZ12, MDZ12. MDZ3 is encoded at chromosome 7422.1, MDZ4 is encoded at chromosome 6501.3-22.2, MDZ7 is encoded at chromosome 16011.2 and MDZ12 is encoded at chromosome 1601.2 and MDZ12 is encoded at chromosome 1601.2 and MDZ12 is encoded at chromosome 1601.2 and MDZ12 is encoded at chromosome 1602.2 associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ1, or MDZ12, e.g. cancer or developmental disorders. The nucleic acused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12. The probes are also be useful as therapautic genetic locus. The probes are protein are useful as therapeutic agents for gene therapy or as protein are useful as therapeutic agents for gene therapy or as protein are useful as therapeutic agents for gene therapy or as

BP; 3 A; 11 C; 5 G; 6 T; 0 U; 0 Other; Sequence 25

Gaps

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Indels

Length 25;

DB 1;

4.3%; Score 18.2; Local Similarity 87.0%; Pred. No. 64; es 20; Conservative 0; Mismatches

Query Match

Matches

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363 TICCICACTITICCIGGACCGCGA 385 2 Trecreacrarecrececes

AAH45474 standard; DNA; 18 BP.

AAH45474/c

RESULT 29

(first entry)

07-SEP-2001

AAH45474;

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This invention relates to a novel method for the detection, treatment and/ or prevention of cellular debilitations or derangements caused by the development of sporadic basal cell carcinoma (BCC). Specifically, it refers to the identification of relevant therapeutic agents based on their effect on the expression level and activity of the Glil transcription factor gene. Glil is a proto-oncogene that is ectopically expressed in epidermal tissue and is linked to tumour formation and neoplasia. The present invention describes cytostatic Glil inhibitors that are useful for detecting the onset or presence of sporadic BCC in an animal. Furthermore, it includes methods for testing the ability of a animal. Furthermore, it includes methods for testing the ability of a drug or other entity to modulate the activity of Glil. This oligonucleotide sequence is the RT-PCR primer Shh-D used to amplify human Shh (secreted sonic hedgehog) RNA of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Upstream activating sequence; transgenic animal; regulatory DNA sequence; hedgehog gene; bigenic animal; transcriptional activating sequence; disease model; cancer; altered vascularisation; brain size regulation; autoimmune disease; tissue proliferation; Parkinson's disease; Shh;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Preventing or treating sporadic basal cell carcinoma by administering an inhibitor of glioma transcription factor-1 (Gli1) activity or expression, and diagnosis of the disease by detecting the presence and level of expression of Gli1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                               RT-PCR; primer; Shh-D; human; ss; PCR; cellular debilitation; sporadic basal cell carcinoma; BCC; Glil; proto-oncogene; tumour formation; neoplasia; cytostatic; secreted sonic hedgehog.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4.2%; Score 18; DB 1; Length 18; 100.0%; Pred. No. 33; 0; Indels ive 0; Mismatches 0; Indels
                                                                                         RT-PCR primer Shh-D used to amplify human Shh RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 18 BP; 4 A; 5 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 6; 22pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16
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ID AAZ49111 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                   97US-0050286P.
98US-00102491.
                                                                                                                                                                                                                                                                                                                                                                         03-APR-2001; 2001US-00825155.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 100.0
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                            (first entry)
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                                                                                                                                                                                                                                                                               US2003100032-A1.
                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                      20-JUN-1997;
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                                          15-JAN-2004
                                                                                                                                                                                                                                                                                                                         29-MAY-2003
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ADD15351;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention relates to a method of detecting the onset or presence of sporadic basal cell carcinoma (BCC) in an animal. The method involves measuring the level of Gili in a sample of Skin. Gili levels above basal or normal indicate the presence or onset of sporadic basal cell carcinoma. Gili is a zinc finger transcription factor down stream of secreted sonic hedgehog (shh) activation in a cascade of cytoplasmic signal transduction. Gili in turn can induce Shh expression in an autoregulatory manner. There are links between ectopic expression of the Gili gene and the development or onset of BCC. The method is useful for detecting the onset or presence of sporadic basal cell carcinoma, particularly in detecting skin cancer. The present sequence represents a PCR primer specific for human Shh cDNA. The primer is used in the method
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Detecting the onset or presence of skin cancer, particularly sporadic basal cell carcinoma, comprises measuring the level of Glil in the

Disclosure, Col 8; 21pp; English.

sample.

Sporadic basal cell carcinoma, BCC, detection, Gli1, skin cancer, transcription factor, PCR primer, human, 88; sonic hedgehog; 8hh. PCR primer Shh-D specific for human secreted sonic hedgehog cDNA.

Homo sapiens,

JS6238876-B1

98US-00102491.

22-JUN-1998;

39-MAY-2001

20-JUN-1997;

(UYNY ) UNIV NEW YORK STATE.

WPI; 2001-366473/38.

Altaba ARI;

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Gaps

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4.2%; Score 18; DB 1; Length 18; larity 100.0%; Pred. No. 33; Conservative 0; Mismatches 0; Indels

Similarity

Query Match Best Local Simi Matches 18;

16

GGAGTCTCTGCACTACGA

GGAGTCTCTGCACTACGA 1

18

8

ADD15351/c ADD15351/c LD ADD15351 standard; DNA; 18 BP. XX

RESULT 30

Sequence 18 BP; 4 A; 5 C; 5 G; 4 T; 0 U; 0 Other;

the invention

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This sequence represents a PCR primer for the mouse Shh gene. The invention relates to a transgenic non-human animal (A) whose cells contain a non-viral regulatory DNA sequence (I) (e.g. an upstream a cativating sequence) linked to a recombinant hedgehog gene (II), which was introduced into the mammal, or its ancestor, at an embryonic stage. Bigenic animals (A'), derived from (A) by introducing a transcriptional activating sequence (TAS), are useful as models of disease, particularly cancer (of breast, skin, protate, kidney, lung, or central nervous system, also primitive neuroectodermal tumours and medulloblastoma). CC target genes on signalling pathways involving hedgehog proteins (HP) (e.g. altered vascularisation, regulation of brain size, density and cellular concentration etc.), and for assaying for a temporal requirement of Path in disease progression (particularly of cancers and autoimmune corrector medials can be used to screen for potential therapeutic disease). The animals can be used to screen for potential therapeutic care returned to the subject, specifically for treatment of Parkinson's or proliferation and differentiation. Hedgehog proteins can also be used to expand a population of neural stem cells from a subject, then the cells come a simple cross since the transcription activate otherwise silent transgenes in progeny from a simple cross since the transcription activator and the silent transgene are maintained in separate mouse lines, and abnormal expression is only induced in the bigenic animal. This eliminates the need for microinjection and genotypic screening for each experiment, and many procession expenses or province contained for experiment, and many procession expenses or province contained for experiment, and many procession expenses or province contained for experiment and ex
Alzheimer's disease; spinal cord injury; therapy; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Transgenic animals useful as disease models, e.g. for cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4.2%; Score 17.8; DB 1; Length 21; 90.5%; Pred. No. 52;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     bigenic embryos can be produced by cross-breeding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 21 BP; 5 A; 6 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pred. No. 52;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 20; 44pp; English.
                                                                                                                                                                                                                                                               98US-0087899P.
                                                                                                                                                                                                         99WO-US012417.
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les 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                    Rowitch DH, Mcmahon AP;
                                                                                                                                                                                                                                                                                                              HARD ) HARVARD COLLEGE
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                                                                                                    WO9963052-A2
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                                                                                                                                                                                                                                                               03-JUN-1998;
                                                                                                                                                    39-DEC-1999
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Matches
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Cell comprising exogenous nucleic acid inducing tyrosine hydroxylase expression useful for treating catecholamine-related diseases such as Parkinson's disease, manic depression and schizophrenia.

Example 1; Page 20; 68pp; English

(SALK ) SALK INST BIOLOGICAL STUDIES.

Palmer T,

Sakurada K,

WPI; 2000-656165/63.

21-MAR-2000; 2000WO-US007544

05-0CT-2000

26-MAR-1999;

Rattus norvegicus. WO200058451-A1

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                                                                                                                                                                                                                                                                                    The present invention describes the rat Nurrl coding and protein sequences. The Nurrl protein is involved in the induction of tyrosine shydroxylase expression in adult rat-derived hippocampal progenitor cells. The Nurrl gene and protein can be used in the treatment of catecholaminerlated diseases such as Parkinson's disease, manic depression and schizophrenia. They can also be used to induce tyrosine hydroxylase expression and identify tyrosine hydroxylase related deficiencies, which are linked to the same diseases. The present sequence is a PCR primer used in a method to differentiate adult neural progenitor cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MD23; MD24; MD27; MD212; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                             4.2%; Score 17.8; DB 1; Length 21; 90.5%; Pred. No. 52; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 2 A; 7 C; 5 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human MDZ3 scanning oligonucleotide SEQ ID 1904.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    104 TGACCGCGACCGCAGCAGTA 124
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-AUG-2001; 2001US-00922181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gu Y, Nguyen C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADB00918 standard; DNA; 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19; Conservative
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EP1281758-A2.
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Gaps

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2; Indels

57 GAGGAGTCTCTGCACTACGAG 77

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21 caccacrcracacrardae 1

AAA95383 standard; DNA; 21 BP

RESULT 32 AAA95383, Rat; Nurr1; tyrosine hydroxylase; catecholamine-related disease; Parkinson's disease; manic depression; schizophrenia; PCR primer; ss.

Rat Shh coding sequence PCR primer #2.

(first entry)

12-FEB-2001

AAA95383;

rng.res

New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer. Example 8; SEQ ID NO 1904; 103pp; English.

The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MD23, MD24, MD212, MD212. MD23 is canced at chromosome 7422.1, MD24 is encoded at chromosome 7422.1, MD27 is encoded at chromosome 6921.3-222.2, MD27 is encoded at chromosome 1691.2 and MD212 is encoded at chromosome 1601.2 and MD212 is encoded at chromosome 500.1 in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MD23, MD24, MD27, or MD212, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MD23, MD24, MD27, or MD212. The nucleic acids can also be used as probes to detect and characterize gross alterations in MD23, MD24, MD27, or MD212 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as protein are useful as therapeutic agents for gene therapy or as

Sequence 25 BP; 3 A; 12 C; 4 G; 6 T; 0 U; 0 Other;

ô Ouery Match 4.1%; Score 17.6; DB 1; Length 25; Best Local Similarity 83.3%; Pred. No. 85; Matches 20; Conservative 0; Mismatches 4; Indels

ò 셤 RESULT 34 ABS55991/c ID ABS55991 standard; DNA; 22 BP.

ABS55991; THE STANK BY SECTION OF SECTION O

23-JAN-2003 (first entry)

Mouse RT-PCR primer Shh rp #1.

Mouse; primer, ss; Hedgehog signalling pathway; T-cell mediated disease; T-cell apoptosis; Notch signalling pathway; cancer; breast; prostate; ovary; T-cell activation; T-cell proliferation; lymphoma; carcinoma; autoimmune disease; inflammatory disease; proliferative disorder; viral infection; genetic immunodeficiency; neurodegenerative disease; myelodysplastic syndrome; isohaemic injury; toxin-induced disease; wasting disease; RT-PCR; reverse transcriptase; Shh; sonic hedgehog.

Mus musculus,

WO200280952-A2

17-OCT-2002.

09-APR-2002; 2002WO-GB001666.

09-APR-2001; 2001GB-00008872. 09-APR-2001; 2001GB-00008873.

(LORA-) LORANTIS LTD

Lamb JR, Hoyne GF, Dallman MJ, Champion BR;

WPI; 2003-058470/05

Use of a modulator of Hedgehog signaling pathways for treating T-cell mediated disease or infection and diseases associated with increased or decreased T-cell apoptosis and T-cell proliferation.

The invention relates to use of a modulator of a Hedgehog signalling pathway or a modulator of a target of the pathway in the preparation of a medicament for treating T-cell mediated disease or infection or a disease or disorder associated with increased or decreased T-cell apoptosis and for modulation of the Notch signalling pathway in cell apoptosis, and for modulation of the Notch signalling pathway in cimume cells. The modulation is useful for treating cancer of the breast, prostate or ovary, lymphomas and carcinomas, autoimmune diseases such as systemic lupus erythematosus, multiple sclerosis and diseases such as systemic lupus erythematosus, multiple sclerosis and diseases such as oppositions such as obtendarthitis and Crohn's diseases such as oppositions such as AlbS and herpesviruses, genetic immunodeficiencies, neurodegenerative diseases such as Alzhaimer's disease and Parkinson's disease, myelodysplastic syndromes such as aplastic anaemia, isochaemic cirrhosis and wasting diseases such as aplastic anaemia, isochaemic cirrhosis and wasting diseases such as cachexia. This sequence represents a reverse transcriptase PCR (RT-PCR) primer used in the scope of the 10; Page 110; 154pp; English.

Sequence 22 BP; 6 A; 5 C; 6 G; 5 T; 0 U; 0 Other;

invention

Gaps .; 0 4.0%; Score 17.2; DB 1; Length 22; 86.4%; Pred. No. 76; tive 0; Mismatches 3; Indels Local Similarity 86.4 les 19, Conservative Query Match Best Loca Matches

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RESULT 35 ADB00922

ADB00922 standard; DNA; 25

ADB00922;

20-NOV-2003 (first entry)

Human MDZ3 scanning oligonucleotide SEQ ID 1908.

Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MD23; MD24; MD27; MD212; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.

Homo sapiens.

EP1281758-A2.

05-FEB-2003.

30-JUL-2002; 2002EP-00016874.

02-AUG-2001; 2001US-00922181

(AEOM-) AEOMICA INC.

ີ່ Shannon M, Gu Y, Nguyen

WPI; 2003-423107/40.

New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer. 

Example 8; SEQ ID NO 1908; 103pp; English.

The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is

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encoded at chromosome 7q22.1, MDZ4 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 1sp1.2 and MDZ12 is encoded at chromosome 1sp1.2 and MDZ12 is encoded at chromosome 1sp26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease acids or MDZ3, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.

Sequence 25 BP; 3 A; 11 C; 4 G; 7 T; 0 U; 0 Other;

Match 4.0%; Score 17.2; DB 1; Length 25; Local Similarity 86.4%; Pred. No. 1e+02; les 19; Conservative 0; Mismatches 3; Indels 364 rccrcacrrrccrggaccgcga 385 Query Match Best Local Si Matches 19 ò

1 rccrcacrarccracccccccc 셤

ACK14726 standard; DNA; 25 BP. RESULT 36 ACK14726 

ACK14726;

14-OCT-2003 (first entry)

Human microarray DNA oligonucleotide SEQ ID NO 114707.

expressed sequence tag; microarray; gene expression; on; biallelic marker; polymorphism; human; cross-species comparison. EST; ss; probe; ex genetic variation;

Homo sapiens,

US2003104410-A1.

05-JUN-2003.

15-MAR-2002; 2002US-00098263

16-MAR-2001; 2001US-0276759P.

(AFFY-) AFFYMETRIX INC.

Mittmann MP;

WPI; 2003-567953/53.

New array of nucleic acid probes, useful for in situ hybridization, in Southern, Northern or dot-blot hybridization to identify or detect the sequence or specific mutations of any gene.

Claim 1; SEQ ID NO 114707; 9pp; English.

The invention discloses a microarray comprising a plurality of nucleic acid probes including one of 2,018,500 fully defined sequences, or its perfect mismatch, antisense match or antisense mismatch. Also disclosed is a method of gene expression analysis. The array is used in analysis of genetic variation or in hybridisation to a DNA library, in analysis of genetic variation or in hybridisation of tag-labelled compounds. The nucleic acid probes are specifically designed for analysis of at least one target sequence. The method of analysis comprises hybridising at least one or more nucleic acids to at least two or more hubbidising at least one as one or more nucleic acids to at least two or more probes are attached to a solid support. The analysis comprises monitoring gene expression levels, identifying biallelic markers or polymorphisms,

or family members of a gene and a cross-species comparison. Each of the nucleic acids further comprises a tag sequence. The array of nucleic acid probes is useful in in situ hybridisation, in Southern, Northern or dotblot hybridisation to identify or detect the sequence or specific mutations of any gene, in mapping the 5' termini of mRNA molecules by primer extensions or in screening CDNA or genomic libraries or subclones for additional subclones containing segments of DNA that have been isolated and previously sequenced. The sequence presented is one of the nucleic acid probes incorporated in the microarray. Note: The sequence data for this patent can also be obtained in electronic format directly from USPTO at sequence.html

Sequence 25 BP; 5 A; 9 C; 6 G; 5 T; 0 U; 0 Other;

ö 4.0%; Score 17.2; DB 1; Length 25; 86.4%; Pred. No. 1e+02; 3; Indels 0; Mismatches 19; Conservative Similarity Query Match Best Local S Matches

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Gaps

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RESULT 37 AAA15463

BP. AAA15463 standard; DNA; 25

AAA15463;

(first entry) 21-SEP-2000 PCR primer for a rat connective tissue growth factor DNA.

Rat, connective tissue growth factor; CTGF; cell proliferative disorder; connective tissue cell; scleroderma; arthritis; cirrhosis; hepatic fibrosis; renal fibrosis; atherosclerosis; cardiac fibrosis; adhesion; surgical scarring; PCR primer; ss.

Rattus sp.

WO200027868-A2.

18-MAY-2000.

99WO-US026189. 05-NOV-1999; 98US-00187478. 99US-00292036. 06-NOV-1998;

14-APR-1999;

(FIBR-) FIBROGEN INC.

Sverdrup F, Carmichael DF; Schmidt BF, Allen ML,

WPI; 2000-376484/32.

New rat connective tissue growth factor, its related gene and antisense sequences useful for modulating CTGF and treatment of cell proliferative disorders.

Example 1; Page 37; 55pp; English.

PCR primers AAA15463-64 were used to amplify DNA encoding a rat connective tissue growth factor (CTGF) polypeptide. The polypeptide may play a signaficant role in the normal development, growth and repair of mammalian tissue. Attisance sequences can be used to inhibit the expression of CTGF in a cell. In particular, the antisense sequences are useful for ameliorating cell proliferative disorders associated with CTGF, e.g. overgrowth of cells, e.g. connective tissue cells. The regulation of CTGF activity comprises down-regulation. The disorders, which can be treated, are chosen from scleroderma, arthritis, cirrhosis, hepatic fibrosis, renal fibrosis, atherosclerosis, cardiac fibrosis, adhesions and surgical scarring. The amisense sequences can also be used to detect expression of CTGF in a sample

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PCR primers AAZ99759-60 were used to amplify a 558 bp of the connective tissue growth factor (CTGF) gene. The specification describes methods for treating or preventing fibrosis or a renal disorder associated with overproduction of extracellular matrix, by administering to a subject an agent that modulates, regulates, or inhibits the expression or activity of CTGF, Healthy individuals demonstrate consistently low levels of urinary CTGF, while in patients with kidney disease the mean level of CTGF increased 4-fold. In those patients with diabetes, but as yet undiagnosed kidney disease, a similar increase was seen. The methods and agents are useful for diagnosing, treating or preventing fibrosis, diabetes, hypertension or a renal disorder associated with overproduction of extracellular matrix.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Connective tissue growth factor; CTGF; fibrosis; renal disorder; extracellular matrix; kidney disease; diabetes; hypertension; FCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       primer F used to amplify a 558 bp fragment of the CTGF gene.
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                                                            Length 25;
                                                                                                                        5; Indels
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Seguence 25 BP; 5 A; 4 C; 12 G; 4 T; 0 U; 0 Other;
                                                        4.0%; Score 17; DB 1; B0.0%; Pred. No. 1.1e+02; iive 0; Mismatches 5
                                                                                                                                                                                        162 GACTGGGTGTACTACGAGTCCAAGG 186
                                                                                                                                                                                                                             1 GAGTGGGTGTGACGAGCCCAAGG 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (FIBR-) FIBROGEN INC.
(FORD-) FORD HEALTH SYSTEM HENRY.
                                                                                                                                                                                                                                                                                                                                                                                                              AAZ99759 standard; DNA; 25 BP
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98US-0112855P.
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                                                                                                                     20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      tissue growth factor
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                                                                                          Similarity
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16-DEC-1998;
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                                                            Query Match
Best Local (
                                                                                             Best Loca
Matches
                                                                                                                                                                                                                                                                                                                                                     RESULT 38
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New array of nucleic acid probes, useful for in situ hybridization, in Southern, Northern or dot-blot hybridization to identify or detect the sequence or specific mutations of any gene.

Claim 1; SEQ ID NO 66407; 9pp; English.

EST; ss; probe; expressed sequence tag; microarray; gene expression; genetic variation; biallelic marker; polymorphism; human;

cross-species comparison

US2003104410-A1. Homo sapiens.

05-JUN-2003.

15-MAR-2002; 2002US-00098263 16-MAR-2001; 2001US-0276759P

(AFFY-) AFFYMETRIX INC.

WPI; 2003-567953/53.

Mittmann MP;

Human microarray DNA oligonucleotide SEQ ID NO 66407.

(first entry)

14-OCT-2003

ACI66416;

ACI66416 standard; DNA; 25 BP.

RESULT 39 ACI66416

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The invention discloses a microarray comprising a plurality of nucleic acid probes including one of 2,018,500 fully defined sequences, or its compered match, perfect mismatch, antisense match or antisense mismatch. Also disclosed is a method of gene expression analysis. The array is used in analysis of genetic variation or in hybridisation to a DNA library, and an monitoring gene expression levels by hybridisation of tag-labelled compounds. The nucleic acid probes are specifically designed for analysis compounds. The nucleic acid probes are specifically designed for marked compounds. The nucleic acid probes and detecting the hybridisation. The nucleic acid probes are attached to a solid support. The analysis comprises monitoring compenses are attached to a solid support. The analysis comprises monitoring compenses are attached to a solid support. The analysis comprises monitoring compenses is useful in in a situ hybridisation. The array of nucleic acid further comprises a tag sequence. The array of nucleic acid probes in mapping the 5' termini of mRNA molecules by confidentian of any gene, in mapping the 5' termini of mRNA molecules by confident acidentify or detect the sequence or specific confidentian subclones containing segments of DNA that have been concluded and previously sequenced. The sequence presented is one of the nucleic acid probes incorporated in the microarray. Note: The sequence containing the microarray. Note: The sequence contains the microarray.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 25 BP; 7 A; 9 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 386 CGACGCCCAAGAAGGTCTTCTAC 410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 CGACGACACCAACTAGGTCTTCGAC 25
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Best Local Similarity 80.0
Matches 20; Conservative
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Gaps

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162 GACTGGGTGTACTACGAGTCCAAGG 186

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Query Match Best Local Si Matches 20,

1 GAGTGTGTGACGAGCCCAAGG 25

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Human microarray DNA oligonucleotide SEQ ID NO 8430
   ACI08439 standard; DNA; 25 BP.
            13-OCT-2003 (first entry)
        ACI08439;
RESULT 40
  ACI08439
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expressed sequence tag; microarray; gene expression; on; biallelic marker; polymorphism; human; EST; ss; probe; expressed genetic variation; bialle cross-species comparison.

Homo sapiens

US2003104410-A1.

05-JUN-2003.

15-MAR-2002; 2002US-00098263

16-MAR-2001; 2001US-0276759P.

(AFFY-) AFFYMETRIX INC.

WPI; 2003-567953/53.

Mittmann MP;

New array of nucleic acid probes, useful for in situ hybridization, in Southern, Northern or dot-blot hybridization to identify or detect the sequence or specific mutations of any gene.

Claim 1; SEQ ID NO 8430; 9pp; English.

The invention discloses a microarray comprising a plurality of nucleic acid probes including one of 2,018 500 fully defined sequences, or its perfect match, perfect mismatch, antisense match or antisense mismatch. Also disclosed is a method of gene expression analysis. The array is used in analysis of genetic variation or in bybridisation to a DNA library, compounds. The nucleic acid probes are specifically designed for analysis of at least one target sequence. The method of analysis comprises of at least one or more nucleic acid probes and detecting the hybridisation. The nucleic acid probes are attached to a solid support. The analysis comprises monitoring probes are attached to a solid support. The analysis comprises monitoring compression levels, identifying biallelic markers or polymorphisms, or family members of a gene and a cross-species comparison. Each of the nucleic acids further comprises a tag sequence. The array of nucleic acid probes in situ hybridisation, in Southern, Northern or dotonois and setting the sequence or specific confort hybridisation to identify or detect the sequence or specific confort additional subclones containing segments of DNA that have been isolated and previously sequenced. The sequence presented is one of the nucleic acid probes incorporated in the microarray. Note: The sequence cate for additional subclones containing segments of DNA that have been solated for this patent can also be obtained in electronic format directly conforts and stream of the electronic format directly conforts and stream of the electronic format directly conforts and stream of the electronic format directly conforts and sequence.

Sequence 25 BP; 5 A; 8 C; 9 G; 3 T; 0 U; 0 Other;

Gaps . 0 4.0%; Score 17; DB 1; Length 25; 80.0%; Pred. No. 1.1e+02; Live 0; Mismatches 5; Indels Similarity 80.0 20; Conservative Query Match Best Local

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GACCGCGACGACGCCCAAGAAGG 402 378

à 셤 RESULT 41 AAF27037

BP. AAF27037 standard; DNA; 37

AAF27037;

(first entry) 30-MAR-2001 Human Sonic hedgehog (Shh) mutagenic primer, SEQ ID NO:41.

Sonic hedgehog; Shh; polymer conjugate; polyalkene glycol group; bioavailbility; formulation; neurological disorder; inflammatory disorder; autoimmune disorder; cancer; neurodegenerative disorder; Parkinson's disease; Huntington's disease. Alzhedmer's disease; eneurological injury; stroke; multiple sclerosis; multipnant glioma; medulloblastoma; neuroectodermal tumour; 

Homo sapiens

Synthetic.

WO200073337-A1.

07-DEC-2000.

26-MAY-2000; 2000WO-US014741

99US-0137011P. 99US-0149016P. 01-JUN-1999; 13-AUG-1999;

(BIOJ ) BIOGEN INC.

Garber E; Taylor F, Pepinsky RB,

WPI; 2001-049927/06.

Modified hedgehog protein, useful in the treatment of Parkinson's disease and Huntington's chorea, comprises a polymer containing a polyalkylene glycol group linked to any residue other than the N-terminal and lysine residues

Example 6; Page 77; 157pp; English.

The invention relates to novel polymer conjugates of hedgehog proteins are which have increased bioavailability. The hedgehog proteins are conjugated to a non-naturally-occurring polymer complying a polyalkylene glycol group, with the proviso that the polymer is not conjugated to the glycol group, with the proviso that the polymer is not conjugated to the glycol group, with the proviso that the polymer is not conjugated to the protein used in the conjugate may be a wild-type or mutant sonic hedgehog protein. The hedgehog (thh) or besert hedgehog (thh) protein. The hedgehog (thh) or besert hedgehog (bh) protein or may be a hedgehog fusion protein. The invention also relates to methods of a faint and or type of modification have on the activity of the conjugate may be used in the management of the position and/or type of modification have on the activity of the protein. The hedgehog polymer conjugates may be used in the management of various medical conditions including various neurological disorders.

Commany be used to prevent preventing or ameliorate neurodogenerative of disorders (e.g., Parkinson's disease, mutingon's disease, Alzheimer's disease); age-associated neurological disease; neurological injury and crauma; immunological diseases of the nervous system (e.g., multiple collesses); age-associated neurological disease; neurological injury and concertosish; stroke, and malignant gliomas, medillobiastomas and concertosish; stroke, and malignant gliomas, medillobiastomas and concertosish in increased half-life, altered tissue distribution (such as an introper or sequence of the headehog protein from proteclytic con improved ability to stay in the vasculature for longer periods of the vasculature for longer periods of the headehold muton, or reduced immunogenicity. In particular, the ability to create the headehold management of the headehold protein of the invention to cross the blood-brain barrier, and an increased thermal stability would be an advantage when formulating the hedgehog protein in powder form. The present sequence represents a human sonic hedgehog mutagenic primer used in an exemplification of the

Sequence 37 BP; 6 A; 10 C; 12 G; 9 T; 0 U; 0 Other;

Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc fingar protein; MDZ3; MDZ1; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.

30-JUL-2002; 2002EP-00016874. 02-AUG-2001; 2001US-00922181.

EP1281758-A2.

05-FEB-2003.

Homo sapiens

Nguyen C;

Gu Y,

Shannon M,

WPI; 2003-423107/40.

(AEOM-) AEOMICA INC.

Human MDZ3 scanning oligonucleotide SEQ ID 1903

(first entry)

20-NOV-2003

ADB00917;

BP

25

ADB00917 standard; DNA;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to the use of an active agent stimulating the expression and/or function of serum response factor (SRF). SRF variants and/or members of the SRF signal transduction pathway in eukaryotic cells for the preparation of a therapeutic drug or a pharmaceutical composition for the preparation of a therapeutic drug or a pharmaceutical composition tumour metastasis, auto-immune diseases, disturbances of would healing, lymphocyte homing and disturbances of immune defense mechanisms that are linked with SRF- related cellular malfunctions. Pharmaceutical compositions of the invention are used in treating diseases associated with expression or misexpression of SRF target gene, which include formation of diseases like metastatic cancer which is influenced by the gene UPA-R, diseases like chronic renal failure, cancer and various hypodyycaemias. The present sequence is Shh specific reverse transcription PCR (RT-PCR) primer used in the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Use of active agent stimulating expression of serum response factor, its variants or components of signal transduction pathway of factor in eukaryotic cells, for treating disturbances or illness e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                          tumour invasion; tumour metastasis; auto-immune disease; would healing; lymphocyte homing; immune defense mechanism; chronic renal failure; cellular malfunction; metastatic cancer; illness; hypoglycaemia; RT-PCR; Shh; reverse transcription PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                            Serum response factor; SRF modulator; signal transduction; disturbance;
                                           Gaps
                                           ô
      Length 37;
                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 23 BP; 4 A; 3 C; 8 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 16.6; DB 1;
Pred. No. 1.1e+02;
0; Mismatches 4;
      4.0%; Score 17; DB 1; Le
69.7%; Pred. No. 2.6e+02;
iive 0; Mismatches 10;
                                                                                TGCTGGCCCGCCTGGCGGTGGAGGCCGGCTTCG 162
                                                                                                                     recagagacrecrecagagregecearerres 37
                                                                                                                                                                                                                                                                                                                                  specific reverse RT-PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 7; 58pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            08-SEP-2000; 2000EP-00119741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-SEP-2000; 2000EP-00119741
                                                                                                                                                                                                 AAD34565/c
ID AAD34565 standard; DNA; 23
                                                                                                                                                                                                                                                                                           16-JUL-2002 (first entry)
Query Match
Best Local Similarity 69.7°
Matches 23, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-271068/32
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unidentified
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                                                                                                                                                                                                                                                        AAD34565;
                                                                                                                                                                              RESULT 42
                                                                                                                                                                                                                                                                                                                                     Shh
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                                                                                                                     В
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New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.

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The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MD24, MD24, MD212, MD212. MD23 is canceded at chromosome 7422.1, MD24 is encoded at chromosome 7621.1, MD24 is encoded at chromosome 6721.2, MD27 is encoded at chromosome 16011.2 and MD212 is encoded at chromosome 16011.2 and MD212 is encoded at chromosome 5921.3-22.2, MD27 is encoded at chromosome 16011.2 and MD212 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with MD21, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease acused by altered expression of MD23, MD24, MD27, or MD212. The nucleic acids can also be used as probes to detect and characterize gross alterations in MD23, MD24, MD27, or MD212, genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as proteins. The present sequence was used to illustrate the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3.9%; Score 16.6; DB 1; Length 25; 32.6%; Pred. No. 1.3e+02; Ive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 25 BP; 3 A; 12 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 8; SEQ ID NO 1903; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               361 ACTICCICACTTICCTGGACCGC 383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Agriccicaciarccidececes 25
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les 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ACI14729 standard; DNA; 25
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RESULT 44
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177 GAGTCCAAGGCACATATCCACTG 199

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3.9%; 82.6%;

Query Match
Best Local Similarity 82.6
Matches 19; Conservative

GAATCCAAAGCTCACATCCACTG 1

23

Length 23; 4; Indels

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New array of nucleic acid probes, useful for in situ hybridization, in Southern, Northern or dot-blot hybridization to identify or detect the sequence or specific mutations of any gene.
                        EST; ss; probe; expressed sequence tag; microarray; gene expression; genetic variation; biallelic marker; polymorphism; human; cross-species comparison.
       Human microarray DNA oligonucleotide SEQ ID NO 14720.
                                                                                                                                                                                                                                       Claim 1; SEQ ID NO 14720; 9pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          251 GGGCTCGGCCACGGTGCACCTGG 273
                                                                                                              .5-MAR-2002; 2002US-00098263.
                                                                                                                               6-MAR-2001; 2001US-0276759P
                                                                                                                                                (AFFY-) AFFYMETRIX INC
                                                                                                                                                                                   WPI; 2003-567953/53.
                                                                            US2003104410-A1
                                                            Homo sapiens.
                                                                                                                                                                  Mittmann MP;
                                                                                             05-JUN-2003
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The invention discloses a microarray comprising a plurality of nucleic acid probes including one of 2,018,500 fully defined sequences, or its perfect match, perfect missues mentiones missued.

Also disclosed is a method of gene expression analysis. The array is used in monitoring gene expression levels by hybridisation to a DNA library.

In analysis of genetic variation or in hybridisation of tag-labelled compounds. The nucleic acid probes are specifically designed for analysis of at least one target sequence. The method of analysis comprises of at least one target sequence. The method of analysis comprises of hybridising at least one or more nucleic acid to hybridisation. The nucleic acid probes and detecting the hybridisation. The nucleic acid probes and detecting the hybridisation. The nucleic acid probes are attached to a solid support. The analysis comprises monitoring probes are attached to a solid support. The analysis comprises monitoring probes are attached to a solid support. The analysis of nucleic acid probes is useful in in situ hybridisation, in Southern, Northern or dotucleic acid further comprises a tag sequence. The array of nucleic acid probes is useful in n situ hybridisation, in Southern, Northern or dotuction of any gene, in mapping the 5' termini of mRNA molecules by correct or additional subclones containing sequence presented is one of the nucleic acid probes incorporated in the microarray. Note: The sequence adara for this patent can also be obtained in electronic format directly correctly in the microarray. Note: The sequence.
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ö Gaps .. 0 Query Match
3.9%; Score 16.6; DB 1; Length 25;
Best Local Similarity 82.6%; Pred. No. 1.3e+02;
Matches 19; Conservative 0; Mismatches 4; Indels Sequence 25 BP; 5 A; 9 C; 8 G; 3 T; 0 U; 0 Other;

Gercreeccaceerreacerce 1 23

엄

ACH53354 standard; DNA; 25 ACH53354; ACH53354/ 8X4X6X8

DNA target sequence #2490 useful in array for genetic analyses

(first entry)

16-OCT-2003

PCR primer P24 to convert human antibody CAT-212 to IgG format

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The present invention relates to nucleic acid sequences that are complementary to particular genes, and can be used as probes for a variety of analyses such as gene expression analysis. Each probe comprises 9 or more consecutive nuclectides from at least one of 14936 nucleotide sequences defined in the patent, or their perfect sense match, sense mismatch, antisense match or antisense mismatch oligonucleotides. The probes may be used in a array comprising at least 10 distinct nucleic acid probes. The array is useful in monitoring gene expression nucleic acid probes. The array is useful in monitoring gene expression cariations, and in hybridisation to a DNA library, in analysing genetic variations, and in hybridisations of a gene. The probes are also useful in in situ hybridisations, in screening cDNA or genomic libraries (or derived subclones) for additional clones containing segments of DNA that have been previously isolated and sequenced, in Southern, northern, or dot-blot hybridisation of genomic DNA to identify or detect the sequence of any gene or detect specific mutations in any gene, and in mapping the 5' termini of many molecules by primer strensions. The sequence of the invention are also useful as PCR primers complementary to particular genes with a wide range of analytical uses. CMH50865-ACH65260 represent the target sequences of the invention. Note: The sequence data for this patent was obtained in electronic format directly from the USPTO web site at seqdata.uspto.gov/psipplDEDIETRY.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New probe array useful e.g. for monitoring gene expression levels, for analyzing genetic variations, or for hybridizing tag-labeled compounds, comprises multiple nucleic acid probes.
               Gene expression analysis; array; hybridisation; genetic variation; tag-labelled compound; gene family; in situ hybridisation; library screening; Southern hybridisation; northern hybridisation; dot-blot hybridisation; gene sequence; mtation detection; target sequence; probe; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3.9%; Score 16.6; DB 1; Length 25; 82.6%; Pred. No. 1.38+02; tive 0; Mismatches 4; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; SEQ ID NO 2490; 9pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    167 GGTGTACTACGAGTCCAAGGCAC 189
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                                                                                                                                                                                                                                                                            08-AUG-2002; 2002US-00215112
                                                                                                                                                                                                                                                                                                                     08-AUG-2001; 2001US-0311040P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAD18152 standard; DNA; 21
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-576608/54
                                                                                                                                                                                                                                                                                                                                                             (MITT/) MITTMANN M.
                                                                                                                                                                                        US2003082596-A1
                                                                                                                                               Unidentified
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                                                                                                                                                                                                                                   01-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                         Mittmann M;
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AAD18152
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The invention relates to a specific binding member which binds to human ectaxin. The binding member comprises an antibody variable heavy (VH) domain from CAT-212 (WH) Variable heavy (CDRs). Variable light (VI) domain from CAT-212 (WH) domain and a VH/VI complementary determining regions (CDRs). Botaxin is a chemoattractant protein that binds to a specific receptor which is expressed predominantly on eosinophils. The binding member is useful for neutralising eotaxin, which is useful in treating at ham, eczema and other atopic diseases such as rhinitis, food allergy, conjunctivitis, allergic colitis which are recognised as eosinophilmentary bowel disease which includes eosinophilic colitis/enteritis/spulman's syndrome, vasculitis including Hugues-Stovin gastroenteritis/shulman's syndrome; vasculitis including Hugues-Stovin syndrome, Churg-Starauss syndrome. The present sequence is a PCR primer used for converting encoding human antibody CAT-212 (ScFv-single chain variable region fragment) to IgG DNA (whole antibody) format
Human, eotaxin, CAT-212; antibody, heavy chain variable region, VH; eczema; asthma; atopic disease; dermatological; rhinitis; food allergy, vasotropic; conjunctivitis; allergic colitis; psoriasis; pemphigoid; eosinophil-mediated disease; cellulitis; drug eruption; vasculitis; inflammatory bowel disease; gastroenteritis; PCR primer; se.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human antibodies against eotaxin useful for treating asthma, eczema and other atopic diseases, comprises an antibody variable heavy or variable light domain from CAT-212 or from complementary determining regions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 3 A; 4 C; 11 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CAMB-) CAMBRIDGE ANTIBODY TECHNOLOGY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 11; Page 103; 107pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Smith S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-MAR-2001; 2001WO-GB000927.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-MAR-2000; 2000US-0187246P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Vaughan TJ, Wilton AJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-589944/66.
                                                                                                                                                                                                                                                                                                                                                                                                              WO200166754-A1.
                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-SEP-2001
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                  Gaps
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0
DB 1; Length 21;
                 1; Indels
Score 16.4; DE
Pred. No. 99;
0; Mismatches
                 ö
 3.8%;
         94.48;
        Local Similarity 94.4
Query Match
                 Matches
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g ઠે

ABZ58547 standard; DNA; 22 BP ABZ58547; ABZ58547 

primer X2R for detection of Pragile E site. <u>Б</u>

(first entry)

13-MAY-2003

Fragile E site; diagnosis; microcapillary electrophoresis; human; trinucleotide repeat; screening; PCR; primer; 88.

Homo sapiens.

WO2003014396-A1

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The present invention relates to a method for diagnosis of a multiplication disease of repeated trinucleoride sequence. The methods multiplication disease of repeated trinucleoride sequence by PCR, analysis of the amplified product on microcapilary electrophoresis (CE), analysis of the amplified product. In Fragile E site (FRAXE), in genetic region Xq28, a CCG trinucleoride is repeated 6-25 times in describing the amplified product. In Fragile E site (FRAXE), in genetic region Xq28, a CCG trinucleoride is repeated 6-25 times in healthy subjects and over 200 times in affected individuals. The present sequence is that of reverse primer XXE which is specific to the FRAXE repeated trinucleoride sequence region. It is used with forward primer XZF (see ABZ5546) to detect FRAXE. A diagnosis kit comprising these produced. Use of CE, especially fabricated as an on-chip analysis system, allows the size of the PCR product to be measured rapidly, with accuracy and reproducibility. The method allows diagnosis before the disease develops and determination of whether a silent carrier will develop the
                                                                                                                                                                                                                                                                          Diagnosing multiplication disease of repeated trinucleotide sequences e.g. Huntington's disease, by amplifying repeated trinucleotide sequence region, migrating and separating product by microcapillary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 22 BP; 4 A; 7 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                        Han S;
                                                                                                                                                                                                                                                                                                                                                                                      Claim 12; Page 8; 45pp; English.
                                                                                                                                                                                        Kim H,
                                                              06-AUG-2002; 2002WO-KR001489
                                                                                                     06-AUG-2001; 2001KR-00047301
                                                                                                                                                                                          Baik S,
                                                                                                                                                (BIOM-) BIOMEDLAB CORP
                                                                                                                                                                                                                                 WPI; 2003-256603/25
                                                                                                                                                                                                                                                                                                                                               electrophoresis.
                                                                                                                                                                                          Lee Y,
                    20-FEB-2003.
                                                                                                                                                                                        Kim J,
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Gaps ö 3.8%; Score 16.2; DB 1; Length 22; 85.7%; Pred. No. 1.2e+02; tive 0; Mismatches 3; Indels Similarity 85.7 18; Conservative Query Match Local Matches ઠે

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86 N 66 CTGCACTACGAGGGCCGCGCA 22 ctrcceraceáceccececa

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BP. ADC58126 standard; cDNA; 24 (first entry) 18-DEC-2003 ADC58126;

cancer; PCR; Mastocyte-specific guanine trinucleotidase 17.49 primer mastocyte-specific guanine trinucleotidase; 17.49; HIV; Unidentified 88 primer;

18-APR-2001; 2001CN-00112607 18-APR-2001; 2001CN-00112607 27-NOV-2002. CN1381569-A. 

(BIOW-) BIOWINDOW GENE DEV INC SHANGHAI.

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Matches
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The present invention provides sequences of antisense oligonucleotides targeted at the murine and human B7-1 and B7-2 coding and mRNA sequences. The antisense sequences have phosphorothicate backbones and some nucleotides are 2'-methoxyethoxy residues. The sequences can be used in the treatment of inflammatory and autoimmune disorders, including asthma, invenile diabetes mellitus, myasthenia gravis, Graves' disease, rheumatoid arthritis, allograft rejection, inflammatory bowel disease, multiple sclerosis, psoriaals, systemic lupus erythematosus, contact dermatitis, rhinitis, allergies and cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel compound for diagnosing, preventing and treating immune disorders, comprising an oligonucleotide that specifically hybridizes with a nucleic acid sequence encoding B7 protein.
(systemic) lupus erythematosus, multiple sclerosis, contact dermatitis, rhinitis, allergy, cancer and metastases. The oligonucleotides may also be used to manipulate T cell activation ex vivo; to determine or detect B7 protein expression; for diagnosis; as assay and purification reagents and to study physiological roles of B7 proteins
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human, mouse, B7-1; B7-2; antisense; PCR primer; inflammation; autoimmune disorder; phosphorothioate backbone; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
3.7%; Score 15.8; DB 1; Length 20;
Best Local Similarity 89.5%; Pred. No. 1.2e+02;
Matches 17; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                     3.7%; Score 15.8; DB 1; Length 20; 89.5%; Pred. No. 1.2e+02; rive 0; Mismatches 2; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human B7-1 mRNA antisense oligonucleotide SEQ ID NO: 26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 5 A; 8 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                 Sequence 20 BP; 5 A; 8 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Karras JG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sxample 1; Page 45; 162pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      398 GAAGGICTICTACGIGATC 416
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                                                                                                                                                                                                                                                                                                                                                                                                                   BP
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                                                                                                                                                                                                                                                                                                                                                                                                   :829/c
AAF32829 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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                                                                                                                                                                                                 Local Similarity 89.5
Les 17, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAF32829;
                                                                                                                                                                               Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ss; human; B7; T cell; inflammation; autoimmune disease; cell activation;
                                                                                                                                                                                                                      The invention relates to a novel mastocyte-specific guanine trinucleotidase 17.49. The protein is useful for treating diseases such as cancer and HIV infection. The current sequence represents a primer related to the mastocyte-specific guanine trinucleotidase 17.49 protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New oligo:nucleotide(s) that modulate expression of B7 proteins - used for, e.g. controlling activation and proliferation of T cells, particularly for treatment, diagnosis and prevention of inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                              Polypeptide-mastocyte-specific guanine trinucleotidase-17.49 and polynucleotide for coding it.
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                                                                                                                                                                                                                                                                                                                                                                                                Length 24;
                                                                                                                                                                                                                                                                                                                                                                                            3.8%; Score 16; DB 1; Length 24;
79.2%; Pred. No. 1.6e+02;
ive 0; Mismatches 5; Indels

    .20
    /*tag= a
    /note= "Phosphorothioate linkages"

                                                                                                                                                                                                                                                                                                                                                         Sequence 24 BP; 3 A; 7 C; 9 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human B7-1 targetted oligonucleotide 13801
                                                                                                                                                                               Example 3; SEQ ID NO 3; 32pp; Chinese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GAGGGCGCGCAGTGGACATCACC 98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24 dagederecedeacacacarerec 1
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AAV47987 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity 79.2
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                                                                       WPI; 2003-249033/25.
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modified_base
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                             Mao Y,
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Gaps

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Human; calreticulin; antisense compound; hyperproliferative disorder; cancer; autoimmune disease; viral infection; cardiovascular disease; antisense therapy; cytostatic; immunosuppressive; virucide; antisense; phosphorothioate backbone; ss.
                                                        Human calreticulin antisense oligonucleotide, ISIS 109305.
                                                                                                                                                                                                                                                                     *tag= c
mod_base= OTHER
note= "2'methoxyethyl nucleotides"
                                                                                                                                                                                               "mod_base= OTHER
note= "2'methoxyethyl nucleotides"
                                                                                                                                                               mod_base= OTHER
note= "Phosphorothicate backbone"
                                                                                                                                      ocation/Qualifiers
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/mod_base= m5c
6. .20
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/mod_base= m5c
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mod_base= m5c
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mod_base= m5c
                                                                                                                                                                                                                       *tag= d
mod_base= m5c
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mod_base= m5c
                                                                                                                                                                                                                                                                                                                                                     *tag= h
mod_base= m5c
AAD39512/c
ID AAD39512 standard; DNA; 20 BP.
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/mod_base= r
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*tag= a
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*tag= b
                                        04-OCT-2002 (first entry)
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                                                                                                                                       Key
modified_base
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                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bennett CF,
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                                                                                                                         Synthetic
                         AAD39512;
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Claim 3; Page 82; 109pp; English.
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The invention relates to antisense compounds, compositions and methods for modulating the expression of calreticulin. The compositions comparise antisense compounds, particularly antisense oligonucleotides, targetted to nucleic acids encoding calreticulin. The antisense compound is useful for inhibiting the expression of calreticulin in human cells or tissues. It is also useful for treating a human having a disease or condition associated with calreticulin, e.g., hyperproliferative disorder e.g. cancer, autoimmune disease, viral infection or cardiovascular disease, by inhibiting expression of calreticulin. It is useful for diagnostics, therapeutics, prophylaxis and as research reagents and kits. It is also used in antisense therapy. The present sequence is an antisense compound targetted to human calreticulin. This sequence is an antisense compound gapmer oligonucleotides

Sequence 20 BP; 2 A; 8 C; 4 G; 6 T; 0 U; 0 Other;

Gaps .. 0 3.7%; Score 15.8; DB 1; Length 20; 89.5%; Pred. No. 1.2e+02; rive 0; Mismatches 2; Indels Local Similarity 89.5 Query Match Best Loc Matches

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Human oligonucleotide sequence. ABZ92967 standard; DNA; 20 (first entry) 17-0CT-2003 ABZ92967; RESULT 52 ABZ92967 ID ABZ9

Human, antisense; lung dysfunction, nasal airway dysfunction, antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antisethmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

23-APR-2002; 2002WO-US013135. WO200285308-A2 Homo sapiens. 31-OCT-2002. 

(EPIG-) RPIGENESIS PHARM INC

24-APR-2001; 2001US-0286137P.

Katz E, Pabalan J, Aguilar S; Li Y, Sandrasagra A, Tang L, Shahabuddin Nyce JW, 1 Miller S,

ä

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 8209; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonuclectide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions,

891. SIIT

wed Apr 41 14:36:41 400%

Novel antisense compound targeted to nucleic acid encoding calreticulin, useful for treating a human having disease or condition associated with calreticulin e.g. cancer, viral infection, autoimmune disease.

Wed Apr 41 12:58:41 2004

junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second acrive agent comprising an entinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiathmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition may have a preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing levels of contining surfacturing bronchodilation, increasing levels of ubiquinone or lung inflammation, lung allergies, or a respiratory disease or condition, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed or specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences 

Seguence 20 BP; 0 A; 8 C; 8 G; 4 T; 0 U; 0 Other;

. 0 3.7%; Score 15.8; DB 1; Length 20; 89.5%; Pred. No. 1.2e+02; rive 0; Mismatches 2; Indels 131 GCTGGCCCGCCTGGCGGTG 149 Local Similarity 89.5 nes 17; Conservative Query Match fatches

2 derecededererere 20 ઠે d d

RESULT 53 ADC65851,

851/c ADC65851 standard; DNA; 20

ВP

18-DEC-2003 (first entry)

ADC65851;

Mouse IGF-beta receptor II targeted antisense oligonucleotide #50.

mouse, antisense oligonucleotide; transforming growth factor beta receptor II; TGF-beta receptor II; typerproliferative disorder; breast cancer; autoimmune disorder; rheumatoid arthritis; 2.0-methoxyethyl gapmer; phosphorothioate backbone; ss; murine.

Mus musculus.

WO2003000656-A2.

03-JAN-2003.

19-JUN-2002; 2002WO-US019665.

21-JUN-2001; 2001US-00888361.

(ISIS-) ISIS PHARM INC.

Murray SF, Wyatt JR;

WPI; 2003-175279/17.

New compound having a sequence targeted to a nucleic acid encoding Transforming growth factor beta-receptor II, useful for preparing a composition for treating hyperproliferative disorder e.g., lung, liver, colon or gastric cancer.

Claim 3; SEQ ID NO 147; 141pp; English.

The invention comprises antisense oligonucleotides that are targeted to the nucleic acid encoding transforming growth factor beta (TGF-beta) receptor II. The antisense oligonucleotides of the invention are useful for treating: hyperproliferative disorders (e.g. breast cancer), or an autoimmune disorder (e.g. rheumatoid arthritis). The present DNA sequence

represents a 2'-O-methoxyethyl gapmer oligonucleotide with a phosphorothioate backbone that is targeted to mouse TGF-beta receptor II. ន្តដ្ឋនូ

Sequence 20 BP; 4 A; 7 C; 5 G; 4 T; 0 U; 0 Other;

ö Query Match 3.7%; Score 15.8; DB 1; Length 20; Best Local Similarity 89.5%; Pred. No. 1.2e+02; Matches 17; Conservative 0; Mismatches 2; Indels

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RESULT 54 ADE27764/

ВЪ 7764/c ADE27764 standard; DNA; 20

ADE27764;

(first entry) 29-JAN-2004 Human B7-1 mRNA targeted oligonucleotide SEQ ID 26.

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Gaps

ss; human; B7-1; inflammatory skin disorder; antisense; psoriasis; contact dermatitis; atopic dermatitis; seborrheic dermatitis; nummular dermatitis; generalised exfoliative dermatitis; eczema; critical costimulatory molecule.

Synthetic. Homo sapiens.

US2003176374-A1.

18-SEP-2003.

09-MAY-2001; 2001US-00851871

31-DEC-1996; 96US-00777266. 04-JUN-1999; 99US-00326186. 25-MAY-2000; 2000WO-US014471.

(VICK/) VICKERS T A. (XARR/) KARRAS J G.

Karras JG; Bennett CF, Vickers TA,

WPI; 2003-863863/80.

Treating an inflammatory skin disorder such as psoriasis comprises topically applying an antisense compound targeted to the nucleic acid encoding human B7 protein.

Example 1; SEQ ID NO 26; 88pp; English.

The invention relates to a method of treating an inflammatory skin disorder in an individual by topically applying an antisense compound targeted to a nucleic acid molecule encoding a human B7 protein. The invention is for treating an inflammatory skin disorder in individual. The skin disorder is psoriasis, contact dermatitis, atopic dermatitis, seborrheic dermatitis, nummular dermatitis, generalised exfoliative dermatitis or eczema. The invention effectively modulates critical costimulatory molecules such as the B7 protein. The present sequence represents a human B7-1 targeted oligonucleotide. 

Sequence 20 BP; 5 A; 8 C; 4 G; 3 T; 0 U; 0 Other;

.. 0 Query Match 3.7%; Score 15.8; DB 1; Length 20; Best Local Similarity 89.5%; Pred. No. 1.2e+02; Matches 17; Conservative 0; Mismatches 2; Indels

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Gaps

398 GAAGGICTICTACGIGAIC 416

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Detecting T-cell activation by measuring the amount of MUC-1 expression useful for diagnosing or treating autoimmune or inflammatory disorders, viral disease or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                 Human; MUC-1; detection; T-cell activation; mucin; antiinflammatory; immunomodulator; antirheumatic; antiarthritic; antiallergic; dermatological; antidabetic; nephrotropic; antithyroid; antianaemic; neuroprotective; hepatotropic; uropathic; ophthalmological; antiviral; cytostatic; autoimmune disorder; inflammatory disorder; viral disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 21; 40pp; English.
                                                                                                                                                                           AAZ91293 standard; DNA; 21 BP
19 GAGGGTCTTCTACGTGAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99WO-US012820.
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                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                 Human MUC-1 PCR primer #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cancer; PCR primer; ss
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                                                                                                                     RESULT 55
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A method has been developed for detecting T-cell activation by evaluating the amount of MUC-1 mucin expression in a T-cell compared to a non-activated control. The method is useful for treating disorders associated with T-cell activation, using an agent (artibody/antagonist) that modulates MUC-1 activity. The T-cell activation associated disorders may be autoimmune or inflammatory disorders (e.g. inflammatory arthritis, portasis, allergies, allergic contact dermatis, rhemmatoris and arthritis, psoriasis, allergies, allergic contact dermatis, ankylosing spondylitis, myasthenia gravis, systemic lupus erythematosus, poblyarteritis nodosas, Goodpastures syndrome, isopathic thrombocytopenic purpura, autoimmune haemolytic anaemia, Grave's disease, rhemmatic fever, permicious anaemia, insulin-resistant diabetes mellitus, bullous pemphigus vulgaris, viral myocarditis (Cocksakie B virus response), autoimmune thyroiditis (Hashimoto's disease), male infertility (autoimmune), sarcoidosis, allergic encephalomyelitis multiple sclerosis, Sjorgens disease, Reiter's disease, Celiac disease, cancer. The present sequence represente a PCR primer for human MUC-1, which is used in an example from the present invention

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Sequence 21 BP; 4 A; 5 C; 5 G; 7 T; 0 U; 0 Other;

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Gaps
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Query Match 3.7%; Score 15.8; DB 1; Length 21; Best Local Similarity 89.5%; Pred. No. 1.3e+02; Matches 17; Conservative 0; Mismatches 2; Indels
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231 AAATCGGGAGGCTGCTTCC 249 ATATCGAGAGGCTGCTTCC 21

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97.

Human IGERB coding sequence PCR primer SEQ ID NO:

(first entry)

15-MAY-2001

0 × 2 × 5 × 6 × 6 ×

AAF92239;

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AAF92239 standard; DNA; 22

AAF92239,

The present sequence is a PCR primer for the human muc-1 mRNA. It was used to amplify the sequence in order to determine the expression pattern of the protein. This showed that MUC-1 is a fimunosuppressor, and its antagonists act to reduce overactive immune responses. Thus, MUC-1 required to treat inflammatory disorders such as rheumatoid arthritis, psoriasis, allergic contect dermatitis and ankylosing spondylitis, autoimmune disorders including myasthenia gravis, systemic lupus erythematosus, polyarteritis nodosa, Goodpastures syndrome, isopathic thrombocytopoenic purpura, autoimmune haemolytic anaemia, Graves' disease, rheumatic fever, pernicious anaemia, insulinersistant myocarditis, autoimmune thyroiditis, male infertility, sarcoidosis, viral myocarditis, autoimmune thyroiditis, male infertility, sarcoidosis, allergic encephalomyelitis, multiple sclerosis, Sjorgens disease, Caliac disease, sympathetic ophthalmia and primary biliary cirrhosis, immune disorders, graft versus host disease and Gaps Use of agent capable of intracellularly inhibiting mucin MUC-1 for inducing T-cell-based immunosuppression and for treating autoimmune disorders, transplant rejection and inflammatory disorders. autoimmune disorder; immune disorder; ö Query Match 3.7%; Score 15.8; DB 1; Length 21; Best Local Similarity 89.5%; Pred. No. 1.3e+02; Matches 17; Conservative 0; Mismatches 2; Indels Sequence 21 BP; 4 A; 5 C; 5 G; 7 T; 0 U; 0 Other; MUC-1; immunosuppression; autoimmune di inflammatory disorder; PCR primer; ss. Example 1; Page 35; 51pp; English. 231 AAATCGGGAGGCTGCTTCC 249 ATATCGAGAGGCTGCTTCC 21 BP. 99WO-US029016 98US-0111973P Agrawal B, Longenecker BM; AAA63180 standard; DNA; 21 (first entry) Human muc-1 PCR primer #2. transplant rejection WPI; 2000-423418/36. (BIOM-) BIOMIRA INC. WO200034468-A2. 11-DEC-1998; Homo sapiens 09-DEC-1999; 06-NOV-2000 15-JUN-2000. AAA63180; RESULT 57 **AAA**63180 g ઠે

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Terada R,
                                              WPI; 2003-332936/31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADA14342;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cksl; ss.
                    Iida S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADA14342
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               셤
    ઠે
                                                                                                                                                                                                                                                                                                                                                                                          The present invention provides the protein and coding sequences of characteristic variants of the human immunoglobulin E receptor beta chain (IGERB). These contain single nucleotide polymorphisms (SNPB) which may be indicative of a predisposition to atopy, allergy, asthma, rhinitis and eczema. Also provided are the sequences of probes and primers for use in identifying the genotype of an individual with regards to the IGERB gene. The IGERB gene is found at human chromosome 11q13. The sequences are all useful in therapeutics. The present sequence was used to isolate
Human, immunoglobulin E receptor beta chain; IGERB; chromosome 11q13; allergy; asthma; rhinitis; eczema; single nucleotide polymorphism; SNP; atopy; probe; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gene construct, genome modification, higher plant, plant, marker gene, homologous recombination, cloning site, T-DNA, plant transformation, monocotyledon, Agrobacterium, gene function analysis, PCR primer, ss.
                                                                                                                                                                                                                                                                                                                    Novel polynucleotide useful for therapeutic purposes, comprises nucleotide polymorphisms in immunoglobulin E receptor beta chain gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3.7%; Score 15.6; DB 1; Length 22; 81.8%; Pred. No. 1.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 22 BP; 2 A; 7 C; 5 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                            Stephens JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31 GCTGGGACGAAGATGGCCACCA 52
                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 77; 88pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GCTAGGACGAGAATGGCCAACA 1
                                                                                                                                                     11-AUG-2000; 2000WO-US022175.
                                                                                                                                                                                                              GENA-) GENAISSANCE PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23-AUG-2002; 2002WO-JP008506.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              8-AUG-2001; 2001JP-00258489
                                                                                                                                                                                   99US-0150423P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACF03722 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (NISB ) JAPAN TOBACCO INC. (SYGN ) SYNGENTA LID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18; Conservative
                                                                                                                                                                                                                                                            Denton RR, Kliem SE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR primer WxR-R3831.
                                                                                                                                                                                                                              NAND/) NANDABALAN K.
                                                                                                                                                                                                                                                                                       WPI; 2001-226623/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       VO2003020940-A1
                                                                                            WO200114588-A1
                                                                                                                                                                                   24-AUG-1999;
                                                                 Homo sapiens,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-SEP-2003
                                                                                                                          01-MAR-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ACF03722;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 58
ACF03722/
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The present invention describes a gene construct (1) for modifying the genome of higher plants by homologous recombination. (1) comprises marker genes and cloning sites between the right bottom sequence (BR) and left bottom sequence (BR) and left in the first cloning sites between the right bottom sequence (BR) and left obttom sequence (BL) originating from T-DMR. Also described: (1) a vector for plant transformation containing any of the constructs, particularly with a first cloning site for integration into the 5' region in the homologous recombination of the target gene into the 3' region in the homologous recombination of the target gene into the 3' region in the homologous recombination of the target gene into the host genome, and a second cloning site for integration into the host genome; and (2) producing a genome-modified higher plant (especially a monocotyledon) by using homologous recombination through negative or calluees with the Agrobacterium; (11) infecting plant cells, tissues or calluees produced by homologous recombination through negative or positive selection; (iv) culturing selected cells or tissues into calluees; (v) culturing in callue. regenerating medium to grow into heterozygously modified plants; and (vi) producing homozygously modified plants; and (vi) producing homozygously modified plants; and (vi) producing homozygously modified plants; and the callue of the constructs are useful for modifying the genome of higher plants by homologous recombination without altering the original locus, for the analysis of genomination which is used in an example from the present invention
                                                                                                                                                                                                               A gene construct for modifying the genome of higher plants by homologous recombination without altering the original locus, comprises marker genes and cloning sites between the right and left bottom sequences from T-DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cancer; anti-cks1; antisense oligonucleotide; benign lesion; papilloma; atherosclerosis; psoriasis; autoimmune disease; bacterial infection; viral infection; HIV; hepatitis; herpes; polythemia; mastocytosis; cks1 inhibitor; skp2 inhibitor; cytostatic; antisense therapy; sarcoma; leukaemia; Hodgkin's lymphoma; non-Hodgkin's lymphoma; adenoma; melanoma; carcinoma; colon cancer; pancreatic cancer; cervical cancer; human; Skp2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .
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3.7%; Score 15.6; DB 1; Length 22;
Best Local Similarity 81.8%; Pred. No. 1.6e+02;
Matches. 18; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 22 BP; 2 A; 5 C; 9 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antisense oligonucleotide SEQ ID NO:40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           372 Trecredacecece 393
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 5; Page 22; 48pp; Japanese.
Inagaki Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP
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Page 37

Wed Apr 21 12:58:21 2004

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Treating cancer, e.g. sarcoma, leukemia, (non-)Hodgkin's lymphoma, adenomas, melanomas, carcinomas, colon cancer, pancreatic cancer or cervical cancer, by employing an anti-cks-1 antisense oligonucleotide.
                                                                                         Disclosure, Page 86, 87pp, English.
12-FEB-2002; 2002US-0356906P.
                                            WPI; 2003-689667/65.
              (CHIR ) CHIRON CORP
                             Walter AO,
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The present invention describes a method for treating cancer comprising using an anti-cksl antisense oligomucleotide. Also described: (1) treating benign lesions (e.g. papillomas, atherosclerosis and psoriasis), autoimmune diseases, bacterial infections, viral infections (e.g. HIV infections, hepatitis or herpes infections), polythemia, or mastocytosis using the cksl antisense oligomucleotide or as skp2 inhibitor; (2) treating cancer using a skp2 inhibitor; (3) cksl inhibitor; (4) a ribozyme, a protein, a polypeptide, an antibody or a small molecule; (4) in solated polymucleotide with a sequence comprising a transcriptional initiation region and a sequence encoding an antisense oligomucleotide; (5) a recombinant vector comprising the polymucleotide; and (6) inhibiting the expression of cksl or skp2 in a mammalian cell. cksl and skp2 antisense therapy, and as Cksl and Skp2 inhibitors. The method is used in antisense therapy, and as Cksl and Skp2 inhibitors. The method is used in antisense melanomas, carcoma, leukaemia, (non-lhodykin's lymphoma, adenomas, melanomas, carcinomas, colon cancer, pancreatic cancer, or cervical cancer. The persent sequence represents an antisense oligomucleotide given in the Sequence Listing of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 23 BP; 8 A; 9 C; 5 G; 1 T; 0 U; 0 Other;
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3:7%; Score 15.6; DB 1; Length 23; 81.8%; Pred. No. 1.8e+02; tive 0; Mismatches 4; Indels
                                                        GCGCAGTGGACATCACCACGTC 103
                                                                                    1 GCGCAGCAGACAAACCACGTC 22
                                18; Conservative
                Local Similarity
                                                            82
    Query Match
                    Best Loca
Matches
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Gapa

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Human RFC40kD gene PCR primer SEQ ID NO: 368
       BP
       ABT03847 standard; DNA; 24
                     (first entry)
                      13-SEP-2002
              ABT03847;
RESULT 60
    ABT03847
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Human; cancer; neoplastic disease; tumour specific marker; cytostatic; transcription factor; PCR; primer; ss.

WO200240716-A2 Homo sapiens. 23-MAY-2002.

16-NOV-2000; 2000US-0249508P. 13-NOV-2001; 2001WO-US043461

(CEMI-) CEMINES PPC

WPI; 2002-537346/57

Palm K;

Determining the presence of neoplastic molecular markers, by identifying the presence of markers in host test sample using array of neoplastic molecular marker specific reagents and analyzing the array of the reagents

21; 41pp; English. Example 1; Page

Shamoon BF

Jefferson AB,

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Reinhard

늉 The present invention relates to a method for determining the presence oneoplastic molecular markers in a host, involving the use of neoplastic molecular marker specific reagents to detect such markers and analysing the array of reagents, allowing the identification of the neoplastic disease present. This can be used to determine the best treatment for cancers, in particular neural cell, lung and prostate tumours. The present sequence is a PCR primer useful for detecting the coding sequences of markers of the invention ####X#X00000000X8

Sequence 24 BP; 4 A; 6 C; 6 G; 8 T; 0 U; 0 Other;

ö Gaps ö Length 24; 4, Indels 3.7%; Score 15.6; DB 1; 31.8%; Pred. No. 1.9e+02; 0; Mismatches Query Match 3.7%; Best Local Similarity 81.8%; Matches 18; Conservative

24 51 CACTCAGAGGAGTCTCTGCACT 72 cagricacadercrerefre

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BP.

ABL54647 standard; DNA; 17

ABL54647 RESULT

Human, p53, p53AIPI, p53-dependent apoptosis-associated, apoptosis; cytostatic, cancer, PCR, primer, ss. Human p53AIPI associated PCR primer SEQ ID NO 20. 02-AUG-2001; 2001WO-JP006666. 03-AUG-2000; 2000JP-00240399 (UYTY ) UNIV TOKYO. (ONCO-) ONCOTHERAPY SCI INC. (first entry) Nakamura Y, Arakawa H; WO200212496-A1. Homo sapiens. 31-MAY-2002 14-FEB-2002. ABL54647;

p53-dependent apoptosis-associated protein and its encoding gene p53AIPI, used for screening apoptosis mediated remedies for cancer and as controllers of apoptosis induction. WPI; 2002-217192/27 

Example 7; Page 40; 121pp; Japanese.

The invention relates to human p53-dependent apoptosis-associated protein, P53AIPI comprising fully defined 806, 777, 2659 nucleotide sequences (ABL54631-ABL54633 respectively) given in the specification and proteins having fully defined 124, 86 and 108 amino acid sequences (ABB08837-ABB08839 respectively) given in the specification. The protein and encoded gene have cytostatic activity, are useful in screening for regulators of apoptosis for subsequent use as cancer treatments. The present sequence is that of the Human p53AIPI associated PCR primer, useful to the invention

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RESULT 62 AAX38484/

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The present invention relates to an expression library comprising synthetic or semi-synthetic nucleic acid sequences, not cloned from an immunized source, where the nucleic acid sequences are darived from mutagenised immunoglobulins that are naturally devoid of light chains. The library is useful for the preparation of antibodies having binding specificity for a target antigen which avoids the need for a donor to have been previously immunized with the target antigen. The recombination of heavy and light chains is avoided, therefore preventing the formation of molecules that are non-functional. The number of hypervariable residues in the binding domain is reduced, allowing a more complete reperior to possible binding variants to be obtained. The present sequence is a PCR primer targeted to anchor regions in llama antibodies. The primers (ApA73754 amplified the framework regions F1, and primer and the control of the primer targeted to anchor regions in llama antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Heavy chain; tetanus; toxin; human; monoclonal; antibody; K4.1; hybridoma; immortalisation; in vivo; xenomice; analysis; immunglobulin; diagnosis; research; therapy; B cell; primer; polymerase chain reaction; amplification; PCR; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Expression library comprising nucleic acids not cloned from an immunized source, derived from immunoglobulins naturally devoid of light chains, use for producing antibodies specific for a target antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The primers (AAA73745 to AAA73754) amplified the framework regions F
F2, F2c, F3 and F4. (Updated on 15-SEP-2003 to standardise OS field)
                                                                                              Llama; primer; expression library; antibody; immunization; anchor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Anti-tetanus toxin human antibody heavy chain cDNA primer MG-24Vi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 15.2; DB 1; Length 20;
Pred. No. 1.5e+02;
                                                       Primer F3 used to amplify part of llama antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 2 A; 7 C; 7 G; 3 T; 0 U; 1 Other;
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85.0%; Pred. No. ...
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Frenken LGJ, Van Der Logt CPE;
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                                                                                                                                                                                                                                                                                                                                99EP-00300351
                                                                                                                                                                                                                                                                                                                                                                   (UNIL.) UNILEVER PLC.
(UNIL.) UNILEVER NV.
(HIND-) HINDUSTAN LEVER LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAT38694 standard; DNA; 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
(revised)
(first entry)
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Best Local Similarity 85.0
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-482910/42.
                                                                                                                                                                                                      WO200043507-A1
                                                                                                                         framework; ss
                                                                                                                                                                                                                                                                                                                                19-JAN-1999;
15-SEP-2003
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                                                                                                                                                                 Lama glama.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      X38552) which are nuclease resistant, and comprises about 3-50 nucleotides complementary to the ribonucleotide reductase gene or the secA gene of a microorganism. The antisense oligonucleotides are used to treat mammalian pathological conditions mediated by microorganisms. The oligonucleotides are particularly useful as antimicrobial agents in crop
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Microorganism inhibitor; antisense; nuclease resistant; treatment; ribonucleotide reductase; secA gene; pathological condition; R1 subunit; antimicrobial agent; crop protection; primer; R2 subunit; ss.
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                                                                                                     Gape
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       invention describes novel antisense oligonucleotides (AAX38301-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New oligonucleotides complementary to RR or SecA genes - useful to inhibit growth of microorganisms.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
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3.6%; Score 15.2; DB 1; Length 20;
Best Local Similarity 85.0%; Pred. No. 1.5e+02;
Matches 17; Conservative 0; Mismatches 3; Indels
                                                           DB 1; Length 17;
                                                                                                   1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 3 A; 8 C; 3 G; 6 T; 0 U; 0 Other;
                      Sequence 17 BP; 2 A; 6 C; 2 G; 7 T; 0 U; 0 Other;
                                                           Score 15.4; DB
Pred. No. 98;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                 E. coli SecA antisense oligonucleotide 40.
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                                                                                                     ö
                                                                                                                                             206 GAAAGCAGAGAACTCGG 222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAA73747 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                            ВÞ
                                                           3.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98WO-CA000666
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  97US-0052160P
                                                                                                                                                                                GAAAGCAGAGAACTTGG
                                                                                                                                                                                                                                                                                          AAX38484 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                         Query Match
Best Local Similarity 94.1:
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1999-120874/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
Escherichia coli.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-JUL-1997;
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                                                                                                                                                                                                                                                                                                                                                                          16-JUN-1999
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RESULT 63
AAA73747/
ID AAA7
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AC AAA7

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The invention relates to modification of the half-life of an antibody.

The method of the invention comprises physically linking an antibody.

The method of the invention comprises physically linking an antibody which contains an Pern receptor binding molety (linge, CR2 and CR3)

domains) to a second Form binding molety, (linge, CR2 and CR3)

CC molecules are protected from degradation by the endosomal FcRb/FcRn

CC receptors, which gives them a relatively extended serum half-life

relative to other serum proteins. The presence of a second FcRn binding

CC moiety further extends the serum half-life of an antibody. By increasing

CC moiety further extends the serum half-life of an antibody by increasing

CC clinical treatments is lowered. This could significantly lower costs for

CC treatment, and lead to less frequent hospital visits as fewer doses are

CC required, thereby increasing the quality of life for patients, and

CC potentially reducing the likelihood of toxicity. The technology can also

CC be adapted to extend the serum half life of other proteins, in addition

CC contibudies. Sequences AAAG6862-A66863 represent PCR primers used in an exemplification of the present invention to amplify CDNA generated from

CN bunan monoclonal antibody poly(A+) mRNA expressed in XenoNice. The PCR

CN products were then cloned into pCRII and sequenced
                                                                                   Modifying antibody half life by linking the antibody to an FcRn binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cytotoxic T-lymphocyte antigen-4; CTLA-4; antibody; immune system; hyperimmunity disorder; autoimmune disease; diabetes; graft rejection; proliferative disorder; cancer; immunodeficient disorder; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 3.6%; Score 15.2; DB 1; Length 23; Best Local Similarity 81.0%; Pred. No. 2.1e+02; Matches 17; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 23 BP; 4 A; 5 C; 10 G; 3 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hanke JH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hanson DC, Neveu MJ, Mueller EE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       263 GGTGCACCTGGAGCAGGCGG 283
                                                                                                                                                       Example 1; Page 47; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3 GGTGCAGCTGGAGCAGTCNGG 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Universal human VH primer MG-30.
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    Foord O;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-DEC-1998; 98US-0113647P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAA46857 standard, DNA; 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-OCT-2000 (first entry)
    Junghans R,
                                              WPI; 2000-224282/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (PFIZ ) PFIZER INC. (ABGE-) ABGENIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200037504-A2
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-JUN-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAA46857;
    Gallo M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antibody contg. fully human variable region specifically reactive with antigen - prepd. by immunisation of non-human animal incapable of producing endogenous immunoglobulin (Ig), but capable of producing human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence is a primer for the PCR amplification of the CDNA encoding the heavy chain of the anti-teranus toxin (TT) human monoclonal antibody (MAD) K4:1, which was secreted by the hybridoma K4:1 and obtained by immortalising B cells from xenomice (containing integrated human DNA from the immunoglobulin locus) immunised with TT. The MAD can be used for analysis, diagnosis, research and therapy, particularly for human therapeutic, and in vivo diagnostic applications
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IgGl; immunoglobulin G; FCRn receptor; FCRb; VH region; heavy chain variable region; serum half life; monoclonal antibody; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       y Match 3.6%; Score 15.2; DB 1; Length 23; Local Similarity 81.0%; Pred. No. 2.1e+02; nes 17; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                         Brenner DG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 23 BP; 4 A; 5 C; 10 G; 3 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                         Klapholz S,
                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               263 GGTGCACCTGGAGCAGGGCGG 283
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 7; Page 28; 64pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Universal human VH primer, MG-30.
                                                                                                                                                                                                                                                                                                                                                                       Kucherlapati R, Jakobovits A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAA06862 standard; DNA; 23 BP.
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                                                                                                                                                                                                                                                                                     95WO-US005500.
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                                                                                                                                                                                                                                                                                                                             (CELL-) CELL GENESYS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1996-497628/49.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (ABGE-) ABGENIX INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200009560-A2.
                                         Key
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                                                                                                                                                                                                                                                                                   28-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-AUG-1998;
                                                                                                                                                     WO9634096-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-FEB-2000,
                                                                                                                                                                                            31-OCT-1996
Synthetic.
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Takobovits A,
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                                                                                                              05-MAY-1997;
                                                                                                                29-SEP-1998;
                                                                            12-AUG-2003
                                                                                                      21-NOV-2002
                                                                        ACD10944;
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The invention relates to a novel chimeric or human monoclonal antibody or its antigen-binding portion that specifically binds to and activates belowing 10040. The anti-CD40 antibody of the invention demonstrates cytostatic, virucide, antibodiezerial, immunostimulant and anti-HIV activities and may be useful for treating a hyperproliferative disorder such as cancer, viral and bacterial infection or genetic, primary or combined immunodeficiency conditions including neutropenia or HIV infection. The anti-CD40 antibodies may also be useful for detecting CD40 in a biological sample in vitro or in vivo, as well as during gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New chimeric or human monoclonal antibody or its antigen-binding portion that specifically binds to and activates human CD40, useful for enhancing an immune response in a human, or treating cancer, HIV, neutropenia or
                                       The invention relates to an antibody that binds to an epidermal growth factor receptor (FGF-r) and exhibits inhibition of Expropriation of EXF-r), the degradation of EXF-r, the EXF induced degradation of EXF-r, vascular endothelial cell growth factor (VEGF) production by tumour cells (by greater than 50%) and endothelial cells of by greater than 40%) and also protects threonine phosphorylation of EXF-r and a 63XD protein. The antibody is inhernalised with EXF-r. The antibody may be used for treating tumours such as lung tumours and colon tumours and for treating inflammation and autoimmune diseases. This sequence represents a PCR primer used to amplify cDNA molecules encoding human EXF-r receptor antibodies of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    anti-CD40 monoclonal antibody, CD40; cytostatic; virucide; antibacterial; immunostimulant; anti-HIV; hyperproliferative; cancer; viral; bacterial infection; immunodeficiency; neutropenia; HIV; gene therapy; human; PCR; primer; ss; universal; VH; MG-30.
                                                                                                                                                                                                                                                                                                                                                                                   3.6%; Score 15.2; DB 1; Length 23; 81.0%; Pred. No. 2.1e+02; tive 0; Mismatches 4; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Feng X;
                                                                                                                                                                                                                                                                                                                                           Sequence 23 BP; 4 A; 5 C; 10 G; 3 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Jia X,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 2; SEQ ID NO 118; 177pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Universal human VH PCR primer MG-30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Corvalan J,
Example 3; Page 17; 100pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           263 GGTGCACCTGGAGCAGGGCGG 283
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GGTGCAGCTGGAGCAGTCNGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADE28495 standard; DNA; 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-JAN-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity 81.0
es 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (PFIZ ) PFIZER PROD INC. (ABGE-) ABGENIX INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-441521/41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        viral infections.
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                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; epidermal growth factor receptor; EGF-r; primer; ss; cytostatic; antiinflammatory; immunosuppressive; tyrosine phosphorylation; EGF-2; eggr-radetion; vascular endothelial cell growth factor; VEGF; tumour; endothelial cell; threonine phosphorylation; autoimmune disease; colon; inflammation; lung; cancer; PCR.
                                                                                                                                                                                                                                        The present sequence represents a PCR primer which is used to amplify a fragment of the gene encoding a heavy chain of human antibodies against cytotoxic T-lymphocyte antigen (CTAA)-4. The specification describes an synthetic antibody which is capable of binding CTAA-4. The antibody is composed of a heavy chain variable region, comprising a modified contiguous sequence from a FRI-FRS sequence encoded by a human VH3-33 family gene. The modifications are contained in CDR1, CDR2 and/or framework regions. The antibodies may be used to inhibit CTAA-4 and down-regulate the immune system to treat hyperimmunity disorders (e.g. autoimmune disease, diabetes and graft rejection) and proliferative disorders (e.g. cancer). CTAA-4 stimulatory agents may be used to upregulate immune system to up-regulate immunodeficient disorders
                                                                                              Novel antibodies capable of binding cytotoxic T-lymphocyte antigen (CTLA) -4 containing specified heavy and light chain sequences, useful for treating, e.g. immune disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sully human monoclonal antibodies that bind to epidermal growth factor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human epidermal growth factor receptor (EGF-r) antibody PCR primer #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3.6%; Score 15.2; DB 1; Length 23; 81.0%; Pred. No. 2.1e+02; ive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 23 BP; 4 A; 5 C; 10 G; 3 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Jia X;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  receptors, useful in cancer therapy.
                                                                                                                                                                                                    Example 2; Page 66; 157pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       263 GGTGCACCTGGAGCAGGCCGG 283
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98US-00162280.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 81.01
These 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    JAKOBOVITS A.
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                                                         WPI; 2000-442647/38.
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GALLO M.
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          Corvalan JR;
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AAT32459;

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missing 5' sequence of CDNA encoding a novel canalicular multispecific organic anion transporter (CMOAT) protein, isolated from a human lambda gtil liver cDNA library. The protein is a new member of the ATP-binding cassette (ABC) transporter family. The ATP dependent cMOAT transporter system mediates hepatobilizry excretion in the liver. cMOAT may be a liver-specific homologue of multidrug resistance-associated protein. The nucleic acids are used to provide cells with cMOAT protein activity.

CMOAT protein activity in cells can be enhanced by increasing the level of glutathione, glucuronide and/or sulphate. Antisense constructs, especially derived from another multidrug resistance (WDN) -related protein, e.g. MDR-1, to the nucleic acids and vectors can be used to decrease the level of cMOAT in a cell. The nucleic acids and proteins can be used especially in diagnosis of Dubin-Johnson disease, Rotor disease or another disease involving cMOAT. The cMOAT gene may also be used as a selectable marker gene. (Updated on 25-MAR-2003 to correct PI field.)
                                                                                                                                                        Canalicular multispecific organic anion transporter protein, rown ATP-binding casette transporter family; ABC transporter, hepatobiliary excretion; multidrug resistance-associated protein, cMOAT protein activity; multidrug resistance-related protein; Dubin-Johnson disease; Rotor disease; PCR primer; ss.
                                                                                                                     used to isolate the missing 5' sequence of rat CMOAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA encoding human and rat canalicular multispecific organic transporter proteins - useful for diagnosis and treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 15; DB 1; Length 23;
Pred. No. 2.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Borst
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 23 BP; 7 A; 8 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bosma PJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 393 GCCAAGAAGGTCTTCTACGTGAT 415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CENT AMSTERDAM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 16; 106pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (INTR-) INTROGENE BV.
(MEDI-) ACAD MEDISCH CENT AMSTERDAN
(HETN-) HET NEDERLANDS KANKER INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Johnson disease and Rotor disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oude Elferink RPJ, Paulusma CC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GCCAAGCAGGTGTTCGTCGTGTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP
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                                                           (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAD25481 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1997-435163/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                     PCR primer 2
                                                                                                                                                                                                                                                                                                                                                                                                                             21-FEB-1997;
                                                                                                                                                                                                                                                                                                                                            WO9731111-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-FEB-1996;
                                                           25-MAR-2003
                                                                                 01-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                    28-AUG-1997.
                                                                                                                                                                                                                                                                                                         Rattus sp.
                                                                                                                                                                                                                                                                                    Synthetic
                       AAT94027;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23
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ID AAD2
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AC AAD2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The calpain large subunit 1 gene located on chromosome 15 codes for a calcium activated neutral protease (CANP3) belonging to the calpain family. Mutations in the gene induce limb-girdle muscular dystrophy (LGMD) 2 disease. The gene, and fragments of it, can be used in the prevention, treatment, diagnosis and detection of a predisposition to CAMD2 disease. Bight primers (AAT32456-63) were used to localise the calpain large subunit 1 gene. The results positioned the gene in a region previously defined as 15q15.1-q21.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human novel Calpain large sub:unit 1 gene encoding a calcium dependent protease - used to develop prods. for the diagnosis and treatment of limb -girdle muscular dystrophy 2 disease.
therapy procedures. The current sequence is that of the human anti-CD40 antibody-related PCR primer of the invention.
                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Calpain, subunit, calcium, protease, mutation, treatment, detection, identification, diagnosis, limg girdle muscular dystrophy, LGMD2, calcium activated neutral protease, CANP; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                            Primer (P94in13) for localisation of calpain large subunit 1 gene.
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                                                                                                Length 23;
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                                                                                                                                           4; Indels
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                                                           Sequence 23 BP; 4 A; 5 C; 10 G; 3 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 21 BP; 1 A; 7 C; 3 G; 10 T; 0 U; 0 Other;
                                                                                              Score 15.2; DB 1;
Pred. No. 2.1e+02;
); Mismatches 4;
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                                                                                                                                                                               263 GGTGCACCTGGAGCAGGCGG 283
                                                                                                                                                                                                       3.5%; Sco.
100.0%; Pre
0;
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                                                                                                                                                                                                                                                                                                                        AAT32459 standard; DNA; 21 BP
                                                                                                  3.6%;
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AAT94027/c
ID AAT94027 standard; cDNA; 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21
                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                            Query Match
Best Local Similarity 81.0
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Beckmann J, Richard I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1996-268611/27.
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Gaps

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AAD25481

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(ORTH ) ORTHO-MCNEIL PHARM INC.
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Benzoxazinone derivative, glucose metabolism; lipid metabolism; NIDDM; PPAR gamma; peroxisome proliferator activated receptor gamma; therapy; non-insulin dependant diabetes mallitus; nephropathy; neuropathy; PCOS, atherosclerosis; retinopathy; polycystic ovary syndrome; hypertension; ischaemia; obesity; heart disease; irritable bowel disorder; stroke; reduced insulin sensitivity; inflammation; cataract; aP2 mRNA; probe; ss.
                                                                                  Probe #18 used in aP2 assay for antagonist.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-MAY-2000; 2000US-0203859P.
11-MAY-2001; 2001US-00853798.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-MAY-2001; 2001WO-US015320.
26-MAR-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200187860-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unidentified.
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Rybczynski PJ; Burris TP,

WPI; 2002-082970/11.

Use of benzoxazinone derivatives for treating a subject suffering from a disorder in glucose and lipid metabolism such as non-insulin dependant diabetes mellitus or obesity.

Example 2; Page 34; 45pp; English.

The invention relates to benzoxazinone derivatives useful as peroxisome proliferator activated receptor (PPAR) gamma modulators. The invention desorbates to pharmaceutical compositions comprising benzoxazinone derivatives and methods for treating the onset of a disorder in glucose and lipid metabolism, preferably a condition of reduced insulin sensitivity such as non-insulin dependant diabetes mellitus (NIDDM), polycystic ovary syndrome (PCOS), hypertension, ischaemia, stroke, heart diseases, irritable bowel disorder, inflammation and cataract. The present sequence is a probe designed to anneal to the apz manh and function in the bDNA manh abade with system. This probe used in the apz assay for antagonist which is used in the exemplification of the

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Sequence 23 BP; 5 A; 6 C; 4 G; 8 T; 0 U; 0 Other;

Gaps Match 3.5%; Score 15; DB 1; Length 23; Local Similarity 78.3%; Pred. No. 2.3e+02; les 18; Conservative 0; Mismatches 5; Indels Query Match

TCTACAGCGACTTCCTCACTTTC 374 rcreckérckértektrekakíre 23 352

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AAI68021 standard; DNA; 23 (first entry) 13-MAR-2002 AAI65021; RESULT 72 AAI 6802 

BP

4h-Benzo(1,4)oxazin-3-one compound; glucose metabolism; lipid metabolism; Perzoxisome proliferator activated receptor; therapy, NIDDM; non-insulin dependant diabetes mellitus; nephropathy; neuropathy; NIDDM; atherosclerosis; retinopathy; polycystic ovary syndrome; hypertension; ischaemia; obesity; heart disease; irritable bowel disorder; cataract; anorectic; nephrotropic; ophthalmological; cytostatic; hypotensive; vasorropic; cerbroprotective; cardiant; antiinflammatory; probe;

Probe #18, used in aP2 assay for antagonist.

(first entry)

12-MAR-2002

AAD24705;

AAD24705 standard; DNA; 23

RESULT 73 AAD24705

ap2 mRNA specific oligonucleotide probe.

Peroxisome proliferator activated receptor gamma; benzoxazinone; NIDDM; non-insulin dependant diabetes mellitus; antidiabetic; anorectic; nephrotropic; ophthalmological; antiarteriosclerotic; cytostatic;

WO200187862-A2

aP2 mRNA; ss. Unidentified.

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The invention provides methods of treating a subject suffering from a condition associated with peroxisome proliferator activated receptor gamma (PpAggamma) activity that involves administering a benzoxazinone compound of a specified formula to the subject. The method is useful for treating and inhibiting in a subject the onset of a condition associated with PpAggamma activity such as a condition of reduced insulin sensitivity, non-insulin dependant diabetes mellitus, obesity, nephropathy, retinopathy, atherosclerosis, polycystic ovary gyndrom, hypertension, ischemia, stroke, heart diseases, irritable bowel disorder, inflammation and cataract. Sequences Aal68004-023 represent oligonucleotide probes specific for ap2 used in ap2 mRNA assay for
                                                                                                                                                                                                                                                                                                                                                                  Use of benzoxazinone derivatives for treating a subject suffering from a condition associated with peroxisome proliferator activated receptor gamma activity e.g. non-insulin dependant diabetes mellitus and obesity.
hypotensive; vasotropic; cerebroprotective; cardiant; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                               Turchi IJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 23;
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                                                                                                                                                                                                                                                                                               Combs DW, Rybczynski PJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 23 BP; 5 A; 6 C; 4 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.5%; Score 15; DB 1; 1
ilarity 78.3%; Pred. No. 2.3e+02;
Conservative 0; Mismarchar
                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 7; Page 29; 46pp; English.
                                                                                                                                                                                                                                                            (ORIH ) ORIHO-MCNEIL PHARM INC
                                                                                                                                                                                                    12-MAY-2000; 2000US-0203861P.
11-MAY-2001; 2001US-00854368.
                                                                                                                                                                   11-MAY-2001; 2001WO-US015377.
                     PPARgamma; probe; ap2; ss.
                                                                                                                                                                                                                                                                                                 Burris TP, Demarest KT,
                                                                                                                                                                                                                                                                                                                                  WPI; 2002-082971/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
les 18; Conserv
                                                                                        WO200187861-A2
                                                                                                                               22-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antagonists
                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Loca
Matches
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Yang XB;

Herzog N,

15-NOV-2001; 2001US-0334887P

(TEXA ) UNIV TEXAS SYSTEM Gorenstein D, Luxon BA,

WPI; 2003-513977/48.

Burris TP, Combs DW, Rybczynski PJ; (ORTH ) ORTHO-MCNEIL PHARM INC 11-MAY-2001; 2001WO-US015383. 12-MAY-2000; 2000US-0203860P. 11-MAY-2001; 2001US-00854302. WPI; 2002-055671/07. 22-NOV-2001. 

The patent discloses 4h-Benzo(1,4) oxazin-3-one compounds which are useful as peroxisome proliferator activated receptor (PPAR) gamma agonists and antagonists. The invention also relates to compositions comprising such compounds and methods for treating or inhibiting the onset of a disorder in glucose and lipid metabolism, preferably a condition of reduced insulin sensitivity, such as non-insulin dependent diabetes mellitus (NIDDM), obesity, atherosclerosis, nephropathy, neuropathy, retinopathy, polycystic ovary syndrome, hypertenaion, ischaemia, stroke, heart diseases, irritable bowel disorder, inflammation and cataract. The present DNA sequence is a probe which is designed to anneal to ap2 mRNA and function in the bDNA mRNA detection system. This probe is used in ap2 assay for antagonist in the exemplification of the invention 0; Gaps Use of 4h-benzo(1,4)oxazin-3-one derivatives for treating a subject suffering from a disorder in glucose and lipid metabolism e.g. non-insulin dependant diabetes mellitus and obesity. Query Match
3.5%; Score 15; DB 1; Length 23;
Best Local Similarity 78.3%; Pred. No. 2.3e+02;
Matches 18; Conservative 0; Mismatches 5; Indels Sequence 23 BP; 5 A; 6 C; 4 G; 8 T; 0 U; 0 Other; Example 38; Page 58; 76pp; English.

352 TCTACAGCGACTTCCTCACTTTC 374 1 TCTGCAGTGACTTCGTCAATTC 23 ò 셤

monitoring biological interaction, modified aptamer; phosphorothioate antagonist; antibacterial; phosphorothioate antagonist; antibacterial; munuosuppressive; antitheumatic; antiathritic; antiinflammatory; cytostatic; anti-HIV; antiartheurotry; antiathritic; antiinflammatory; functional proteomics; nuclear factor kappa-B; NF-kappaB; toxic shock; sepsis; rheumatord archritis; Crônn's disease; inflammatory bowel disease; asbestos lung disease; Hodgkin's disease; prostate cancer; ventilator induced lung injury; cancer; AIDS; human cutaneous T cell lymphoma; lymphoid malignancy; HIV-1 induced adult T-cell leukaemia; atherosclerosis; cytomegalovirus; herpes simplex virus; JCV; SV-40; rhinovirus; influenza; Oligonucleotide duplex Seq ID94 related to biological interactions. ADD43640 standard; DNA; 23 BP. (first entry) 402003050290-A2. 15-JAN-2004 Jnidentified 19-JUN-2003. ADD43640; RESULT 74 ADD43640 

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This invention relates to a novel apparatus for monitoring biological interaction which comprises a substrate and a modified aptamer attached contented with the substrate, where a target molecule or its portion, contacted with the modified aptamer under conditions to allow formation of a complex between the modified aptamer and the target molecule or its portion, is detected. The invention may be useful in developing phosphorotrhioare agonists or antagonists which may have antibacterial, immunosuppressive, antirheumatic, antiarthritic, antiarthritic, antiarthritic, antiarthritic, antiarthritic, antiarthritic, and apparatus of the present invention are useful for monitoring hological interactions and in functional proteomics. As an example, and apparatus of the present invention are useful for monitoring biological interactions and in functional proteomics. As an example, and apparatus of the present invention are useful for monitoring biological interactions and in functional proteomics. As an example, and apparatus of the present related diseases, such as toxic shock, sepsis, rectains NF-kappas aptamer-related diseases, such as toxic shock, sepsis, remainated athirtis, Crohn's disease, generalised inflammatory bowel disease, abbestos lung diseases, Hodgkin's disease, prostate cancer, cell lymphoma, lymphoid malignancies, HTM-1 induced adult T-cell leukaemia, atherosclerosis, cytomegalovirus, herpes simplex virus, JCV, SV-40, rhinovirus, influenza, neurological disorders and lymphomas. The present sequence is that of an oligomucleotide duplex which was used during the exemplification of the invention.
                                                                                                                                                                                                                                                      New apparatus with a substrate and a modified nucleotide aptamer for monitoring biological interactions, useful for diagnosing and treating NF-kB aptamer-related diseases, such as toxic shock, rheumatoid arthritis,
                                                                                                                                                                                                                                                                                                                                                                                                            Claim 58; SEQ ID NO 94; 67pp; English.
                                                                                                                                                                                                                                                                                                                                                       cancer and AIDS.
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Gaps . 3.5%; Score 15; DB 1; Length 23; 78.3%; Pred. No. 2.3e+02; 5; Indels Human G-alpha-13 antisense inhibitor ISIS# 20741. 0; Mismatches 132 CTGGCCCGCCTGGCGGTGGAGGC 154 1 crerrccaecreeceereeeec 23 AAZ31792 standard; DNA; 18 BP (first entry) Query Match 3.57 Best Local Similarity 78.3 Matches 18; Conservative 24-JAN-2000 AAZ31792; RESULT 75 AAZ31792 a a ठ

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Sequence 23 BP; 1 A; 6 C; 11 G; 5 T; 0 U; 0 Other;

G-alpha-13; human; inhibitor; cancer; antisense compound; therapy; ss. 98US-00205860. 98US-00205860. Homo sapiens. 04-DEC-1998; 04-DEC-1998; US5981732-A. 09-NOV-1999 Synthetic 

(ISIS-) ISIS PHARM INC

07-AUG-2002; 2002WO-US025049.

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WPI; 1999-633376/54.
                                                                            Escherichia coli.
                                                                               DE19846499-A1.
                                                                                       09-OCT-1998;
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                                                                                   0-APR-2000
 Cowsert LM
                                                          AAA12163;
                                                     AAA12163
                                                  RESULT
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nucleic acid molecule encoding human mitochondrial phosphoenolpyruvate carboxykinase (also known as PEPCK-mitochondrial phosphoenolpyruvate carboxykinase (also known as PEPCK-mitochondrial) phosphoenolpyruvate mitpEPCK), where the Oligonucleotide specifically hybridise with and inhibit the expression of human mtPEPCK. The antisense oligonucleotides can be used for inhibiting the expression of mtPEPCK in human cells or particularly a human suspected of having or treating an animal, particularly a human suspected of having or being prone to a condition or disease associated with expression of mtPEPCK. They can also be used in diagnostics and as research reagents in sandwich and other assays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compound targeted to a nucleic acid molecule encoding human mitochondrial phosphoenolpyruvate carboxykinase useful for treating a human with a mitochondrial phosphoenolpyruvate carboxykinase-associated
amplify the ilvC gene, panE gene, panB gene, panC gene, panD gene and the avtA gene which are used in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human, mitochondrial phosphoenolpyruvate carboxykinase; PEPCK-M; PCK2; PEPCK-mitochondrial; mtPEPCK; antisense oligonucleotide; modulation; phosphorothioate; inhibition; diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human mtPEPCK phosphorothioate antisense oligonucleotide SEQ ID NO:11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                           Length 20;
                                                                                                                                                       2; Indels

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    /*tag= a
    /note= "phosphorothioate linkages"

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                                                                  Sequence 20 BP; 4 A; 7 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                           Score 14.8; DB 1;
Pred. No. 1.9e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
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                                                                                                                  3.5%;
                                                                                                                                                                                                                                  3 AGTCTCTTCACTACCAGG
                                                                                                                                                                                                  61 AGTCTCTGCACTACGAGG
                                                                                                                                                                                                                                                                                                                                                       AAZ95323 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                       1 Similarity 88.9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cowsert LM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-205209/18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        03-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                            31-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-FEB-2000.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mckay R,
                                                                                                                                                                                                                                                                                                                                                                                                  AAZ95323;
                                                                                                                    Query Match
Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo
                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                RESULT 77
AAZ95323
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention describes a novel method for the production, and improvement, of panthothenic acid (1)-producing microorganisms by amplifying (particularly overexpressing) sequences (1) that encode ketopanthoate reductase (XPR), specifically the pans gene, either individually or together. Optionally the ilvC gene is also amplified. (1) is a vitamin used in cosmetics, medicine and human or animal nutrition. The method provides increased yields of (1), e.g. 35-40 mug/ml for the most productive strains. AAA12160-A12171 represent PCR primers used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               useful as
that
                                                                                                                                                                                                  This sequence represents an antisense inhibitor of the invention, and inhibits the expression of the human G-alpha-13 protein. The antisense compounds of the invention are of 8 to 30 nucleobases in length, that inhibits the expression of the human G-alpha-13. The antisense compound is useful for treating an animal, particularly humans, having or being prone to a disease or condition associated with the expression of G-alpha-13, such as cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PCR primer; panthothenic acid; ketopanthoate reductase; panB gene; vitamin; cosmetic; medicine; nutrition; ilvC gene; panB gene; panC gene; panD gene; avtA gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Production of microorganisms that overproduce pantothenic acid, vitamin in e.g. foods or medicines, by overexpressing sequences encode ketopantothenate reductase.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Dohmen J;
                                                                                                                         Antisense compound inhibiting expression of human G-alpha-13
                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
3.5%; Score 14.8; DB 1; Length 18;
Best Local Similarity 88.9%; Pred. No. 1.5e+02;
Matches 16; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                              Sequence 18 BP; 5 A; 5 C; 8 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   E. coli ilvC gene PCR primer panE2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 4; Page 9; 24pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  122
                                                                                                                                                                    Claim 11; Col 38; 38pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Elischewski F, Kalinowski J,
Farwick M, Thierbach G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GACCGCGACGCCAGGAAG 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAA12163 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-AUG-2000 (first entry)
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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an infilammatory steroid in a subject, for reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine reducing surfactant in a subject's tissue, or treating bronchoconstriction, lung surfactant in a subject's tissue, or treating bronchoconstriction, note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                 Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiatehmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodiation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pabalan J, Aguilar D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 4 A; 6 C; 6 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sandrasagra A, Katz E,
i, Shahabuddin S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 15; SEQ ID NO 509; 872pp; English
TACGGCATGCTGGCCCGC 140
                                   3 TACGGCATGATGGCCAGC 20
                                                                                                                                                                                                                                                                    Human oligonucleotide sequence
                                                                                                                             267/c
ABZ85267 standard; DNA; 20 BP.
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Tang L,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
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Miller S,
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The invention relates to a novel DNA vaccine for flatfish rhabdovirus (HIRRV) infected fishes, which provides immunity against HIRRV. The vaccination method uses a DNA construct comprising a transcriptional-control sequence containing cytemegalovirus immediate-type promoter, operably coupled to a nucleotide sequence encoding an immunogenic polyapeptide of HIRRV. The DNA vaccine has vircide activity. The HIRRV DNA vaccine is useful for administering to a fish belonging to the flatfish family by gene gun. The HIRRV DNA vaccine is useful for inducing preventing HIRRV infected by HIRRV and is also useful for preventing HIRRV infection in flatfish. The HIRRV DNA vaccine is effective in enhancing immunity of fish infected by HIRRV. This effective in enhancing immunity of fish infected by HIRRV This infected by Appreciation is equence represents an oligo used in the analysis of the mRNA expression level from the muscles of flatfish, following an innoculation with the flatfish rhabdovirus vaccine of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA vaccine for flatfish rhabdovirus infected fishes has DNA construct comprising a transcriptional control sequence coupled to a nucleotide sequence encoding an immunogenic protein of flatfish rhabdovirus.
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                                                                                                                                                                                                                                                              DNA vaccine; flatfish rhabdovirus; HIRRV; fish; immunity; transcriptional-control; cytomegalovirus immediate-type promoter; immunogenic; virucide; gene gun; ss; primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 6 A; 4 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  rcagaagacrcrcrarac 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (MEIJ ) MEIJI SEIKA KAISHA LTD (AOKI/) AOKI H.
                                                                                                                                                                                                                                Flatfish rhabdovirus oligo #33
CTGCACTACGAGGGCCGC 83
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                                                                                                                       BP
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10-SEP-2001; 2001JP-00274202.
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AAA49039 standard; DNA; 20
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                                                                                                                                                                                              (first entry)
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                                                                                                                       ADD22542 standard; DNA;
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Best Local Similarity
Matches 16; Conservat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-818526/77.
                                                                                                                                                                                                                                                                                                                                      Hirame rhabdovirus
                                                                                                                                                                                                                                                                                                                                                                            JP2003155254-A.
                                                                                                                                                                                            15-JAN-2004
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99
                                                                                                                                                           ADD22542;
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AAA49039/c
ID AAA49
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AC AAA49
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Gaps

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Query Match
3.5%; Score 14.8; DB 1; Length 20;
Best Local Similarity 88.9%; Pred. No. 1.9e+02;
Matches 16; Conservative 0; Mismatches 2; Indels

Rats FH;

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AAT51423-T51435 represent amplification primers used in the construction of an E. coli hygromycin B phosphotransferase (hpt) gene containing vector of the invention. The vector these sequences were used to construct also contained a luciferase gene. The hpt gene used in the vector, is used as a dominant selectable marker. The hpt gene has preferably been modified, to provide increased resistance to hygromycin in comparison to the wild type gene. In the vector, the hpt gene has the control of a promoter (such as the OpP-2 promoter) that is native to gazicus bisporus. The vector can then be used in the production of a starker, and domor marker using this vector; the selection marker, and domor maker integrated into the homobasidomycetes, and expressed at a level which allows direct selection, and stable maintenance of the transformed cells. Previously, the domor DNA was not both integrated and expressed at high enough levels for direct selection and stable maintenance to be possible. The transformed homobasidiomycetes can then be used for the commercial production of substances, such as
                                                                                                                                                                                                                                                                                         Production of stably transformed homo-basidiomycetes - with altered genetic characteristics for e.g. commercial production of enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21 BP; 6 A; 7 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                      (ATOD-) ATO-DLO INST AGROTECHNOLOGISCH ONDERZOEK.
(CNCC-) CNC COOEPERATIVE NEDERLANDSE CHAMPIGNONK.
                                                                                                                                                                                     Huizing HJ,
                                                                                                                                                                                                                                                                                                                                                                        Claim 37; Page 27; 86pp; English
                                                                                                                                                                                     Mooibroek A, Van De Rhee MD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAZ26124 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human polymorphic region 313.
94WO-NL000164.
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Best Local Similarity 81.0
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          enzymes and metabolites
                                                                                                                                                                                                                                        WPI; 1995-067335/09.
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  13-JUL-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to the Thermus sp. AK16D DNA ligase enzyme. This thermostable ligase has 100 fold higher fidelity than T4 ligase and 6 fold higher fidelity than Thermus thermophilus ligase. The present sequence is the degenerate antisense primer #3 corresponding to amino acids 641-647 of the T.thermophilus HB8 DNA ligase gene. This primer was used to amplify DNA ligase gene fragments from various Thermus sprains. The high specificity and thermostability of Thermus sp. Ak16D ligase makes it useful for use in ligase based linear signal amplification, known as LDR/PCR. Ligation of suitable oligomucleotide probes can be disrupted by hybridisation mismatches. This feature may be used to detect infectious diseases (for example bacterial, fungal or viral infection),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Polymerase chain reaction, PCR; primer; amplify; E. coli; GDP-2 promoter; Agaricus bisporus; hygromycin B phosphotransferase; hpt gene; luciferase; homobasidiomycetes; metabolite; enzyme production; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                               Degenerate primer #3 targeted to T.thermophilus HB8 DNA ligase gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New thermostable DNA ligase for sealing a ligation junction between oligonucleotide probes and the target sequence.
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                                                                                                                            Thermostable ligase; bacterial; fungal; viral; infection; cancer genetic disease; PCR primer; antisense; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 1 A; 6 C; 2 G; 5 T; 0 U; 6 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 2; Page 24; 55pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (CORR ) CORNELL RES FOUND INC.
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                         (first entry)
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                                                                                                                                                                                                               Thermus thermophilus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cao W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Primer Nco-HPTS.
                                                                                                                                                                                                                                                                  WO200026381-A2.
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                                                                                                                                                                                                                                                                                                                                                                        29-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                 30-OCT-1998;
                            10-JAN-2001
                                                                                                                                                                                                                                                                                                                        11-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Barany F,
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Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; premalignant metaplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss.
                             Gaps
                             ö
3.4%; Score 14.6; DB 1; Length 21; 81.0%; Pred. No. 2.3e+02;
                            4; Indels
                             0; Mismatches
                                                          90 GACATCACCACGTCTGACCGC 110
                                                                          1 GACATCACCATGCTGAACTC 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                     98WO-US005419.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 97US-0041057P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (VARI-) VARIAGENICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-MAR-1998;
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WO9502691-A2

Synthetic.

AAT51423;

RESULT

Query Match

Best Loc Matches

282 20

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26-JAN-1995

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AAZ26557 standard; DNA; 21 BP
                                                                                                                                                         Human polymorphic region 746.
                                                                                                                                                   30-NOV-1999 (first entry)
                                                                                                      Local Similarity 81.0 ses 17; Conservative
         WPI; 1998-521232/44.
                                                                                                                                                                                           WO9841648-A2
                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                  24-SEP-1998.
   Housman D,
                                                                                                                                             AAZ26557;
                                                                                                    Query Match
                                                                                                          Matches
                                                                                                                                RESULT
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This invention describes a novel method for identifying an inhibitor of potentially useful for treatment of cancer, where the inhibitor is active on a gene vital for cell growth or viablity, and where the gene is subject to loss of heterozygosity (LOH) in a cancer. The inhibitor is used for preventing the development of cancer in a patient having a conscioun, by administering to the patient a first allele present in cells of the precancerous condition, where the normal somatic present in cells of the precancerous condition, where the normal somatic cells of the patient are heterozygous for the first gene, the inhibitor is active on at least one but less than all allelic forms of the gene comman somatic cells, and the first gene. The producte and methods can be consent in the diagnosis, prevention and treatment of LOH disorders, e.g. cancers, atherosclerotic plaques, premalignant metaplastic or dysplastic cancers, benigh tumours, endometriods. Polyvestic kindry disease, and cancers where host disease. The method can also be used to remove malignant cells from bone marrow transplants. AAZ25812-Z26825 represent con man polymorphic sites described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Respiratory syncytial virus; RSV; viral vector; mutated RSV gene; HBV; RSV antigenome; functional deletion; M2 gene; RSV-A; RSV-B; antigen; L gene mutation; vaccine; bivalent vaccine; influenza; HIV-1; HIV-2; SS.
                   Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.4%; Score 14.6; DB 1; Length 21; 81.0%; Pred; No. 2.3e+02; tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oligonucleotide used to construct recombinant RSV vaccines.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21 BP; 2 A; 8 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    241 GCTGCTTCCCGGGCTCGGCCA 261
                                                                                                                         Disclosure; Fig 7; 605pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21 GCGGCTTCCCAGGCAGGCCA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP.
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98US-0084133P.
98US-0089207P.
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AAX35042 standard; DNA; 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 81.v.
Local 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9915631-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-SEP-1997;
04-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-APR-1999.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
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AAX35042/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             셤
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                                                                                                                                                                                                                                                                         This invention describes a novel method for identifying an inhibitor of potentially useful for treatment of cancer, where the inhibitor is active on a gene vital for cell growth or viablity, and where the gene is subject to loss of heterozygosity (LOH) in a cancer. The inhibitor is used for preventing the development of cancer in a patient having a precancerous condition, by administering to the patient a first allele specific inhibitor (ASI) targeted to an allele of a first essential gene specific inhibitor for the precancerous condition, where the normal somatic cells of the patient are heterozygous for the first gene, the inhibitor is active on at least one but less than all allelic forms of the gene present in a population and targets only one allelic forms of the gene comment of colls of the green in the diagnosis, prevention and treatment of LOH disorders, e.g. cancers, atherosclerotic plaques, premalignant metaplastic or dysplastic concers, benigh tumours, endometriosis, polycystic kidney disease, and graft versus host disease. The method can also be used to remove malignant cells from bone marrow transplants. AZZSSB12-ZZSB25 represent thuman polymorphic sites described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; premalignant metaplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss.
                                                                                                                            Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3.4%; Score 14.6; DB 1; Length 21;
81.0%; Pred. No. 2.3e+02;
ative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 2 A; 12 C; 4 G; 3 T; 0 U; 0 Other;
                              Stanton VP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                326 GGCGGCGACGACCAGGCCG 346
                                                                                                                                                                                                                                     Disclosure, Fig 7; 605pp; English.
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                              Ledley FD,
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Recombinant respiratory syncytial viruses.

Bryant

Tang R, Li S,

Jin H,

WPI; 1999-244413/20

Example 6; Page 35; 85pp; English

Housman D, Ledley FD, Stanton VP;

WPI; 1998-521232/44

VARI-) VARIAGENICS INC

97US-0041057P.

19-MAR-1998; 20-MAR-1997;

Gaps ö Sequence 21 BP; 3 A; 9 C; 7 G; 2 T; 0 U; 0 Other;

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The specification describes recombinant respiratory syncytial virus (RSV) particles and viral vectors which express heterologous genes or mutated RSV genes. The RSV particles comprise a RSV antigenome or genome containing at least one functional deletion in an M2 gene, or encode antigenic polypeptides of both RSV-A and RSV-B, or contain a L gene mutation. The recombinant RSV particles can be used to produce vaccines, e.g. bivalent vaccine against RSV-A and RSV-B, or RSV and influenza. Recombinant RSV vaccines can also be constructed for viruses such as HIV-1, HIV-2 and HBV, by constructing a RSV comprising a heterologous sequence from these organisms. The present oligonucleotide was used to construct the ribozyme/T7 terminator sequence, which was construct vectors which are used in the course of the invention

Sequence 21 BP; 2 A; 7 C; 9 G; 3 T; 0 U; 0 Other;

Gaps ., 0 Query Match 3.4%; Score 14.6; DB 1; Length 21; Best Local Similarity 81.0%; Pred. No. 2.3e+02; Matches 17; Conservative 0; Mismatches 4; Indels

31 GCTGGGACGAAGATGGCCACC 51 21 GCTGGGACCATGCCGGCCACC 1 à g

RESULT 85 AAX35043

AAX35043 standard; DNA; 21 BP. AAX35043; 

01-JUL-1999 (first entry)

Oligonucleotide used to construct recombinant RSV vaccines.

Respiratory syncytial virus; RSV; viral vector; mutated RSV gene; HBV; RSV antigenome; functional deletion; M2 gene; RSV-A; RSV-B; antigen; L gene mutation; vaccine; bivalent vaccine; influenza; HIV-1; HIV-2; se.

Synthetic.

WO9915631-A1

01-APR-1999,

28-SEP-1998;

98WO-US020230.

97US-0060153P. 98US-0084133P. 98US-0089207P. 26-SEP-1997; 04-MAY-1998; 12-JUN-1998;

(AVIR-) AVIRON INC.

Li S, Tang R, Jin H,

WPI; 1999-244413/20.

Recombinant respiratory syncytial viruses.

Example 6; Page 35; 85pp; English.

The specification describes recombinant respiratory syncytial virus (RSV) particles and viral vectors which express heterologous genes or mutated RSV genes. The RSV particles comprise a RSV antigenome or genome containing at least one functional deletion in an M2 gene, or encode antigenic polypeptides of both RSV-A and RSV-B, or contain a L gene mutation. The recombinant RSV particles can be used to produce vaccines, e.g. bivalent vaccine against RSV-A and RSV-B, or RSV and influenza. Recombinant RSV vaccines can also be constructed for viruses such as HIV-1, HIV-2 and HBV, by constructing a RSV comprising a heterologous construct the ribozyme/T7 terminator sequence, which was construct vectors which are used in the course of the invention

RESULT 87 AAZ48457

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Characterizing drug-target interactions and identifying genetic mutations that confer resistance to antibacterial compounds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence is a PCR primer for the coding sequence for enoylacrate (Tabl) from Neissaria gonorrheae. The protein was used to create a number of mutants which can be used to determine the targets of antibacterial compounds and understand how the target and compound interacts. This in turn is useful for identifying other antibacterial agents. The Fabl sequence is particularly useful for generating thy dinylocoxydiphenylether (DHDPE) resistant strains of N. gonorrheae, Staphylocoxous surens, Streptococcus progrems, Pacinical Actinobacter, E. coli, Staphylococcus aurens, Streptococcus progrems, Pseudomonas aeruginosa, Enterococcus faecium, Bacillus subtilis and Halloobacter pylori, which can then be used to determine how to fight infection by these bacteria. This primer was used to create random mutations in the Fabl coding sequence. (Updated on 15-SEP-2003 to
                                                                                                                                                                                                                                                                                                                                                          Fabl; encyl-ACP reductase; DHDPE resistance; infection; PCR primer; ss.
                                     Gaps
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Query Match 3.4%; Score 14.6; DB 1; Length 21; Best Local Similarity 81.0%; Pred. No. 2.3e+02; Matches 17; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21 BP; 5 A; 7 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                      Neisseria gonorrheae Fabi PCR primer Gc8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 23; 55pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 266 GCACCTGGAGCAGGCGGCAC 286
                                                                    31 GCTGGGACGAAGATGGCCACC 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 dcacchecaecardcegrac 21
                                                                                                      1 GCTGGGACCATGCCGGCCACC 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28-OCT-1998; 98US-0105965P.
                                                                                                                                                                                               AAA53282 standard; DNA; 21
                                                                                                                                                                                                                                                                   (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (WARN ) WARNER LAMBERT CO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                            Neisseria gonorrhoeae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Dunham SA, Olson E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2000-350764/30.
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                                                                                                                                                                                                                                                                                                                                                                                                                              WO200024932-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-SEP-1999;
                                                                                                                                                                                                                                                                 15-SEP-2003
05-OCT-2000
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                                                                                                                                                                                                                                 AAA53282;
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88

catecholamine-related disease; sion: schizophrenia; PCR primer;

AAZ48457;

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The present invention describes the rat Nurrl coding and protein sequences. The Nurrl protein is involved in the induction of tyrosine hydroxylase expression in adult rat-derived hippocampal progenitor cells. The Nurrl gene and protein can be used in the treatment of catecholaminerlated diseases such as Parkinson's disease, manic depression and schizophrenia. They can also be used to induce tyrosine hydroxylase expression and identify tyrosine hydroxylase related deficiencies, which are linked to the same diseases. The present sequence is a PCR primer used in a method to differentiate adult neural progenitor cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; variant thrombospondin 1; variant thrombospondin 4; SNP;
polymorphism; vascular disease; coronary artery disease; forensics;
myocardial infarction; atherosclerosis; stroke; venous thromboembolism;
pulmonary embolism; paternity test; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cell comprising exogenous nucleic acid inducing tyrosine hydroxylase expression useful for treating catecholamine-related diseases such as Parkinson's disease, manic depression and schizophrenia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /*tag= a
/standard_name= "single nucleotide polymorphism"
                        hydroxylase; catecholamine-relate manic depression; schizophrenia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 5 A; 4 C; 7 G; 5 T; 0 U; 0 Other;
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Pred. No. 2.3e+02;
0; Mismatches 4;
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replace(11,a)
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                                                                                                                                                                                                                                                                                                                                                                     Gage FH;
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                                                                                                                                                                                                                                                                             99US-00277078.
                                                                                                                                                                                                                            21-MAR-2000; 2000WO-US007544
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Best Local Similarity 81.0%;
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                                                                                                                                                                                                                                                                                                                                                                     Sakurada K, Palmer T,
                        Rat; Nurrl; tyrosine
Parkinson's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-656165/63.
                                                                                          Rattus norvegicus.
                                                                                                                                    WO200058451-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                  05-OCT-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Variation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention provides a novel in vitro method for the detection of microorganisms and viruses. The method comprises: (1) forming a comprises chair reaction (FCR) mixture by combining a predetermined volume of a sample to be tested for the presence of a nucleic acid sequence comprising 5'-TAGAAGC-3', known amounts of a first primer comprising 5'-GCTAAGCTCCCAAAGCT-3', and a second primer comprising 5'-CAGAGCTCCTAACA-3', and PCR reagents; (2) forming a PCR product by cycling the PCR mixture to amplify the nucleic acid sequence, if present, comprising 5'-GGTGGCTCTCTAAGCGCC-3' to replicate and attain 0.25-10000mug nuclectide product kmul mixture; (3) adding a probe containing DNA comprising 5'-GGTGGCTCTTAAGCCAC-3' to the PCR mixture or to the PCR product to cause the DNA to hybridize with concern and expense of sequence, if present, and change the conformation of the probe; and (4) determining whether or not bacteria are present in the sample by detecting the conformational change of the probe, a conformational change indicating the presence of bacteria in the sample. CT the methods can be used for the detection of viruses and microorganisms, including bacteria, yeast, molds and protister. They can be used in the food and cosmetic industry and in clinical diagnostics. Using the method
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Detection of microorganisms and viruses, for use in the food and cosmetic industries and for clinical diagnostics.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  including bacteria, yeast, molds and protista. They can be used in food and cosmetic industry and in clinical diagnostics. Using the of is not necessary to remove non-hybridized probe from the system
                                                                                                                                                                           Microorganism, virus, polymerase chain reaction, food, cosmetic, clinical diagnostic, molecular beacon, PCR primer, ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3.4%; Score 14.6; DB 1; Length 21;
81.0%; Pred. No. 2.3e+02;
iive 0; Mismatches 4; Indels
                                                                                                                               Nucleic acid fragment used in detection of microorganisms.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 5 A; 6 C; 7 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 37; Page 38; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GCTGGGACGAAGATGGCCACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GGTGGCTCGAAGATAGCCACC
                                                                                                                                                                                                                                                                                                                                                                                        99WO-US010940.
                                                                                                                                                                                                                                                                                                                                                                                                                                    98US-0086025P.
21
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ID AAA95400 standard; DNA; 21
                                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (HUNT-) HUNT WESSON INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fraser MS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-086985/07.
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les 17; Conser
                                                                                                                                                                                                                                               Unidentified
                                                                                                                                                                                                                                                                                              WO9963112-A2
                                                                                                                                                                                                                                                                                                                                                                                      18-MAY-1999;
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                                                                                   27-MAR-2000
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Query Match Best Loca Matches AAA95400;

Romick TL,

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Gaps

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Length 21; 4; Indels

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Nucleic acids comprising single nuclectide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and atherosclerosis.
                                                       (WHED ) WHITEHEAD INST BIOMEDICAL RES. (MILL-) MILLENNIUM PHARM INC.
                                                                                                                                                                        Example; Page 207; 242pp; English
                  10-SEP-1999; 99US-0153357P.
26-JUL-2000; 2000US-0220947P.
16-AUG-2000; 2000US-0225724P.
07-SEP-2000; 2000WO-US024503.
                                                                                                       WPI; 2001-226749/23
                                                                                    Lander ES,
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Matches
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The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various bolymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thromboembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification Gaps ö ch 3.4%; Score 14.6; DB 1; Length 21; 1 Similarity 81.0%; Pred. No. 2.3e+02; 17; Conservative 0; Mismatches 4; Indels Sequence 21 BP; 5 A; 7 C; 5 G; 4 T; 0 U; 0 Other; CGGCTGCTCTACAGCGACTTC 365 CGGCAGTTGTAGAGCGACTTC 1 Query Match Best Local Similarity

AAF25449 standard; DNA; 21 (first entry) 15-MAY-2001 

BP.

RSV; RSV strain A2; RSV subgroup A; virus accessory gene; vaccine; ds Oligonucleotide used to construct a ribozyme/T7 terminator sequence. Synthetic.

02-AUG-2000; 2000WO-US021079. WO200108703-A1. 38-FEB-2001.

99US-00368076 (AVIR-) AVIRON 03-AUG-1999;

Bryant M; Tang R, Li S, 4PI; 2001-191424/19 Jin H,

Infectious respiratory syncytial virus particle, useful for producing vaccines, comprises a viral genome or antigenome with a deletion in an

Disclosure; Page 34; 128pp; English.

terminator sequence, which was then ligated to the ends of the CDNA respiratory syncytial virus (RSV). The specification describes an infectious RSV particle comprising an RSV (anti) spenome that has at least one functional deletion in a virus accessory gene. Especially, the genome contains the reverse complement of a mRNA-encoding sequence linked to a polymerase-binding site (BBS) of an RSV. The RSV particles of the invention are useful for preparing attenuated, live vaccines, including those that express heterologous gene products (particularly from another strain of RSV, some other virus or pathogen, cellular protein or tumour strain of RSV, some opter virus or pathogen, cellular protein or tumour strain of RSV, some products (e.g. viral proteins or ribozymes for prevention or treatment of disease) in cells and/or to rescue heterologous genes in virus particles Oligonucleotides AAF25449-59 were used to construct a ribozyme/T7

Mccarthy JJ;

Daley GQ,

Bolk S,

Gargill M, Ireland JS,

Seguence 21 BP; 2 A; 7 C; 9 G; 3 T; 0 U; 0 Other;

Gaps ô Query Match 3.4%; Score 14.6; DB 1; Length 21; Best Local Similarity 81.0%; Pred. No. 2.3e+02; Matches 17; Conservative 0; Mismatches 4; Indels

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31 GCTGGGACGAAGATGGCCACC 51 

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BP ABK96224 standard; DNA; 21 ABK96224/c

ABK96224; 

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Respiratory syncytial virus genome construction oligonucleotide #1. (first entry) 24-SEP-2002

Respiratory syncytial virus; RSV; attenuated phenotype; antigeno G protein; F protein; M2-2 gene; expression vector; vaccine; ss.

Synthetic.

WO200244334-A2. 06-JUN-2002

28-NOV-2000; 2000US-00724416. 28-NOV-2001; 2001WO-US044819.

(AVIR-) AVIRON INC.

Ë à Tang Jin H,

Bryant

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WPI; 2002-508507/54.

Isolated infectious respiratory syncytial virus particle, useful as a vaccine, has an attenuated phenotype comprising the viral genome that has a heterologous sequence encoding a G and F protein and a mutation in the a heterolo M2-2 gene.

Example 6; Page 39; 150pp; English.

The invention describes an isolated infectious respiratory syncytial virus (RSV) particle with an attenuated phenotype comprising an RSV antigenome or genome, where the genome or antigenome has a heterologous sequence encoding a G and F protein, and a mutation in the M2-2 gene. The SKV particle is useful as expression vector or vaccine. This sequence represents an oligonuclectide used in the construction of leader and trailer sequences for creation and functional analysis of reporter

1 CTGATTGACAGGGACTTCCTC 21

g

BP

21

ADC49462 standard;

ADC49462 ID ADC4 RESULT

(first entry)

18-DEC-2003

ADC49462;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR; primer; ss; Fc receptor gamma chain; neuroprotective; oligodendroglia; myelin; neurodegenerative disease; multiple sclerosis; myelin formation disorder; Krabbe's a disease; adrenoleukodystrophy; metachromic leukodystrophy; Fc receptor III alpha gamma chain.
plasmids and construction of a cDNA representing the complete genome of
                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Drug compositions containing Fc receptor gamma chain activator for treatment and prevention of neurodegenerative disorders including multiple sclerosis.
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                                                                                           Match 3.4%; Score 14.6; DB 1; Length 21; Local Similarity 81.0%; Pred. No. 2.3e+02; les 17; Conservative 0; Mismatches 4; Indels
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                                                        Seguence 21 BP; 2 A; 7 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fc receptor III alpha gamma chain PCR primer #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 9; Page 49; 109pp; Japanese.
                                                                                                                                                                                 31 GCTGGGACGAAGATGGCCACC 51
                                                                                                                                                                                                           21 GCTGGGACCATGCCGGCCACC 1
                                                                                                                                                                                                                                                                                                                             ABX95654 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-JUL-2002; 2002WO-JP007378.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-JUL-2001; 2001JP-00229553.
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                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-248118/24.
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                                                                                                                                                                                                                                                                                                                                                                       ABX95654;
                                                                                                Query Match
                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                      RESULT 92
                                                                                                                                                                                                                                                                                                         ABX95654
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                                                                                                   non-human animal model; demyelinating disease; myelinogenesis inhibition; myelinogenesis signal molecules; oligodendroglia; screening; myelin growth regulator; multiple sclerosis; PCR; primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention comprises a non-human animal model for demyelinating disease - in which myelinogenesis is inhibited by a defect of myelinogenesis signal molecules in oligodendroglia. The non-human animal model of the invention is useful for screening for a myelin growth regulator, or for screening for a therapeutic agent which is useful for treating a demyelinating disease such as multiple sclerosis. The present DNA sequence represents a PCR primer that was used in an example of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Multipotent adult stem cell; MASC; cell replacement therapy; cytostatic; cardiant; cardiovascular; hepatotropic; haemostatic; antidiabetic; virucide; antinflammatory; vasotropic; antianaemic; neuroprotective;
                                                                                                                                                                                                                                                                                                                                                                               Novel non-human animal model for demyelinating disease in which wellnogenesis is inhibited by defect of myelinogenesis signal molecules in oligodendroglia, for screening for therapeutic agent for multiple sclerosis.
                                                                       Non-human animal model for demyelinating disease-related PCR primer #10.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21 BP; 4 A; 6 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                       (KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example; SEQ ID NO 10; 56pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          348 CTGCTCTACAGCGACTTCCTC 368
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       crearreacaeceacriccic 21
                                                                                                                                                                                                                                                                                           10-SEP-2001; 2001JP-00274232.
                                                                                                                                                                                                                                                            10-SEP-2001; 2001JP-00274232.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABN84964 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (revised)
                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-630032/60.
                                                                                                                                                                                                  JP2003079270-A.
                                                                                                                                                                   Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-AUG-2003
25-NOV-2002
                                                                                                                                                                                                                               18-MAR-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    invention.
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Gaps

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348 CIGCICTACAGGGACTICCIC 368

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Retrovirus LTR PCR primer.

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The present sequence is a primer for a retrovirus long terminal repeat (LTR). The primer was used in an example from the invention in which a retroviral marking study was used to demonstrate that neurons, astrocytes and oligodendrocytes can produced from a single bone marrow-derived multipotent adult stem (MASC), which also differentiated into muscle and differentiating the invention relates to methods of obtaining, maintaining and differentiating MASC. The MASC are derived from a non-embryonic organ or tissue, such as bone marrow, muscle, brain, umbilical cord blood or placenta, and have the capacity to be induced to differentiate to a cell type of mesodermal, ectodermal or endodermal origin, including of mesole, smooth muscle, cardiac muscle, fibroblast, marrow stroma, skeletal muscle, smooth muscle, cardiac muscle, endothelial, pither, categories, handled the constitutively express oct4 and high levels of telomerase and are negative for CD44, MHC class I and MHC class II expression. Teratomas are not formed when MASC are administered to a patient. MASC or their progeny are particularly useful for treating cancer, cardiacyascular disease, conficiency, connective tissue disorders, autoimmune disease, desenerative or traumatic neurological conditions, autoimmune disease, desenerative or traumatic neurological conditions, autoimmune disease, be directed to abdominal aortic aneurysm, cardiac bypass surgery, peripheral vascular disease, or coronary vascular disease (all claimed). (Updated on 07-AUG-2003 to correct OS field) (Updated on 29-AUG-2003 to correct OS field) (Updated on 29-AUG-2003 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                             New multipotent adult stem cells that can be induced to differentiate to form a cell type of mesodermal, ectodermal or endodermal origin, useful for treating e.g. cancer, diabetes, hepatitis, hemophilia, ischemia or
  cerebroprotective; immunosuppressive; antibacterial; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 10; Page 55; 117pp; English.
                                                                                                                                                                                                                                                   14-FEB-2001; 2001US-0268786F.
15-FEB-2001; 2001US-026906ZP.
7-AUG-2001; 2001US-03166Z5F.
25-OCT-2001; 2001US-0343386F.
                                                                                                                                                                                                         14-FEB-2002; 2002WO-US004652
                                              unidentified retrovirus.
                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-667000/71.
                                                                                                                                                                                                                                                                                                                                                                      (ANON ) ANONYMOUS.
                                                                                                                 WO200264748-A2.
                                                                    Unidentified.
                                                                                                                                                             22-AUG-2002.
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                                                      Query Match

3.4%; Score 14.6; DB 1; Length 22;
Best Local Similarity 81.0%; Pred. No. 2.5e+02;
Matches 17; Conservative 0; Mismatches 4; Indels
Sequence 22 BP; 3 A; 4 C; 8 G; 7 T; 0 U; 0 Other;
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ઠ 원 RESULT 95
ABN84966/C
ID ABN84966 standard; DNA; 22 BP
XX
AC ABN84966;
XX
DY 79-AUG-2003 (revised)
DT 07-AUG-2003 (first entry)

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Multipotent adult stem cell; MASC; cell replacement therapy; cytostatic; cardiant; cardiovascular; hepatotropic; haemostatic; antidiabetic; virucide; antinflammatory; vasotropic; antianaemic; neuroprotective; cerebroprotective; immunosuppressive; antibacterial; PCR; primer; ss.
                                                                                                                 14-FEB-2001; 2001US-0268786P.
15-FEB-2001; 2001US-0269062P.
07-AUG-2001; 2001US-0310625P.
25-OCT-2001; 2001US-0343386P.
                                                                                                    14-FEB-2002; 2002WO-US004652
                                                     unidentified retrovirus.
Unidentified.
                                                                                                                                                     (ANON ) ANONYMOUS.
                                                                          WO200264748-A2
                                                                                        22-AUG-2002.
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New multipotent adult stem cells that can be induced to differentiate to form a cell type of mesodermal, ectodermal or endodermal origin, useful for treating e.g. cancer, diabetes, hepatitis, hemophilia, ischemia or inflammation.

WPI; 2002-667000/71.

Example 10; Page 55; 117pp; English.

The present sequence is a primer for a retrovirus long terminal repeat (ITR). The primer was used in an example from the invention in which a retroviral marking study was used to demonstrate that neurons, astrocytes and oligodendrocytes can produced from a single bone marrow-derived multipotent adult stem (MASC), which also differentiated into muscle and differentiating masc. The MASC are derived from a non-embryonic organ or tissue, such as bone marrow, muscle, brain, umbilical cord blood or placenta, and have the capacity to be induced to differentiate to a cell type of mesodermal, ectodermal or endockmal origin, including corteoblast, chondrocyte, adipocyte, fibroblast, marrow stroma, skeletal corteoblast, chondrocyte, adipocyte, fibroblast, marrow stroma, skeletal muscle, smooth muscle, cardiac muscle, endothelial, pither, corteoblast, chondrocyte cell types.

CC panoreas, haematopoletic, gilal, neuronal or oligodendrocyte cell types. CC mestabolic disease oct4 and high levels of telomerase and are not formed when MASC are administered to a patient. MASC or their progeny are particularly useful for trreating cancer, cardiovascular disease, cort canneation entrological conditions, autoimmune disease, disease, transplant rejection, ischaemia or inflammation. Treatment may be directed to abdominal aortic aneurysm, cardiac bypass surgery, peripheral vascular disease, or coronary vascular disease (all claimed). (Updated on 07-MUG-2003 to correct OS field.) (Updated on 29-AUG-2003 to correct OS field.) (Updated on 29-AUG-2003 to

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Gaps
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                                          Query Match
3.4%; Score 14.6; DB 1; Length 22;
Best Local Similarity 81.0%; Pred. No. 2.5e+02;
Matches 17; Conservative 0; Mismatches 4; Indels
Sequence 22 BP; 3 A; 4 C; 8 G; 7 T; 0 U; 0 Other;
                                                                                                                                                   49 ACCACTCAGAGGAGTCTCTGC 69
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21 Arcacrcadaddadcctrcc 1 AAC72998 standard; DNA; 17 RESULT 96 AAC72998/c ID AAC7299 셤

09-FEB-2001

AAC72998;

WO200058519-A2. Homo sapiens.

05-CCT-2000,

31-MAR-1999;

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qenetic analysis.
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Lipshutz RJ,
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Lipshutz RJ,
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                                                                                                31-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-OCT-2000,
                              05-OCT-2000.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention is concerned with a number of human single mucleotide polymorphisms (SNPs) which the inventors identified in human genes. These SNPs can be used in disease diagnosis and prediction of an individual's susceptibility to disease, in forensic and paternity testing and in genetic mapping. In particular, the SNPs of the invention can be used to disganose usceptibility to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, schizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's
                                                                                                           Single nucleotide polymorphism; SNP; human; genetic disease;
disease susceptibility; cardiovascular system; endocrine system;
neurological system; forensic testing; paternity testing; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Single nucleotide polymorphism; SNP; human; genetic disease;
disease susceptibility; cardiovascular system; endocrine system;
neurological system; forensic testing; paternity testing; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nucleic acid selected from one of 106 genes comprising single nucleotide polymorphisms, allele-specific oligonucleotides to the genes are useful for phenotypic correlations, forensics, paternity testing, medicine and genetic analysis.
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                                                                                                                                                                                                                                                                                                                                                                                                Altshuler D, Cargill M, Daley GQ, Ireland JS, Lander ES;
Lipshutz RJ, Patil N, Sklar P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.4%; Score 14.4; DB 1; Length 17;
93.8%; Pred. No. 1.6e+02;
tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 17 BP; 1 A; 7 C; 6 G; 3 T; 0 U; 0 Other;
                                                                          Single nucleotide polymorphism PCR primer #1885.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Single nucleotide polymorphism PCR primer #1881.
                                                                                                                                                                                                                                                                                                                                               (WHED ) WHITEHEAD INST BIOMEDICAL RES. (AFFY-) AFFYMETRIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 8; Fig 5; 214pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           303 CTGAGCCCCGGGGACC 318
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                                                                                                                                                                                                                                                                              30-MAR-2000; 2000WO-US008440.
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                                           (first entry)
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Best Local Similarity
Matches 15; Conserv
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diseases

Homo sapiens

09-FEB-2001

AAC72992;

AAC72992/c RESULT 97

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The present invention is concerned with a number of human single nucleotide polymorphisms (SNPs) which the inventors identified in human genes. These SNPs can be used in diesase diagnosis and prediction of an individual's susceptibility to disease, in forensic and paternity testing and in genetic mapping. In particular, the SNPs of the invention can be used to diagnose usceptibility to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, schizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nucleic acid selected from one of 106 genes comprising single nucleotide polymorphisms, allele-specific oligonucleotides to the genes are useful for phenotypic correlations, forensics, paternity testing, medicine and
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                                                                                                                                                                                                                                                                      , Daley GQ, Ireland JS, Lander ES;
Sklar P;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 1 A; 7 C; 6 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                      (WHED ) WHITEHEAD INST BIOMEDICAL RES. (APPY-) APPYMETRIX INC.
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30-MAR-2000; 2000WO-US008440
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                                                                                                                                                                                                                                                                      Cargill M,
Patil N, S
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Single nucleotide polymorphism; SNP; human; genetic disease; disease susceptibility; cardiovascular system; endocrine system; neurological system; forensic testing; paternity testing; PCR primer; ss.
                                                                                                                                                            The present invention is concerned with a number of human single nucleotide polymorphisms (SNPs) which the inventors identified in human genes. These SNPs can be used in disease diagnosis and prediction of an individual's susceptibility to disease, in forensic and paternity testing and in genetic mapping. In particular, the SNPs of the invention can be used to diagnose susceptibility to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, schizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleic acid selected from one of 106 genes comprising single nucleotide polymorphisms, allele-specific oligonucleotides to the genes are useful for phenotypic correlations, forensics, paternity testing, medicine and
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                                            Nucleic acid selected from one of 106 genes comprising single nucleotide polymorphisms, allele-specific oligonucleotides to the genes are useful for phenotypic correlations, forensics, paternity testing, medicine and
                                                                                                                                                                                                                                                                                                                                               Seguence 17 BP; 1 A; 7 C; 6 G; 3 T; 0 U; 0 Other;
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Best Local Similarity
              WPI; 2000-611722/58
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                                                                                                    genetic analysis.
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Lipshutz RJ,
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(first entry)

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individual's susceptibility to disease, in forensic and paternity testing and in genetic mapping. In particular, the SNPs of the invention can be used to diagnose usceptibility to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, schizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CDK1, PCNA and Cyclin B1. Representative examples of ribozyme recognition sites are given in AAA82415 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is efficient in
                                                                                                                                                                                                                                                                                                                                                                                                                                    Ribozyme; hairpin; hammerhead; gene therapy; vasotropic; restenosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New hairpin and hammerhead ribozyme for inhibiting restenosis, cleaves RNA encoding a cyclin or cell-cycle dependent kinase other than CDK1, PCNA and Cyclin B1.
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                                                                                                                                    3.4%; Score 14.4; DB 1; Length 17; 93.8%; Pred. No. 1.6e+02; Ative 0; Mismatches 1; Indels
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                                                                                                          Sequence 17 BP; 1 A; 7 C; 6 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                         cdk8 ribozyme binding site #90.
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les 15; Conservative
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Best Local Similarity 93.8
Matches 15; Conservative
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Matches
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Gaps ö

1; Indels

3.4%; Score 14.4; DB 1; 93.8%; Pred. No. 1.6e+02; iive 0; Mismatches 1;

CTGAGCCCCGGGGACC 318

CTGAGACCCGGGGGACC 1

Length 17;

RESULT 101

Lander

, Daley GQ, Ireland JS, Sklar P;

Cargill M, Patil N, S

99US-0127248P

dopaminergic gene, DRDS, susceptibility diagnosis; h aura, depression; anxiety; variant allele detection;

97WO-US014830, 96US-0024399P.

21-AUG-1997;

22-AUG-1996; 17-JAN-1997;

(GLAX ) GLAXO GROUP LTD

WPI; 1998-168887/15.

Peroutka SJ;

gene.

Primer 40DRD5.SB.PCR2 for DRD5

migraine with aura; differentiation; ss

PCR primer;

Homo sapiens.

Synthetic

WO9807426-A1

26-FEB-1998.

(first entry)

09-JUL-1998

AAV25487;

AAV25487 standard; DNA; 20

AAV25487

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The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a ribozyme (1) which cleaves RNA encoding a cytokine involved in inflammation, matrix metalloproteinase (MMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid segment encoding (I). (I) can have antipsoriatic, dermatological, cytostatic, antiseborrheic, antidiabetic, antisickling, ceptablamlological, vulnerary, keratolytic and virucide activities, and cleaves RNA encoding cytokine involved in inflammation. (I) can be used in gene therapy. (I) and (II) are useful for treating proliferative skin diseases such as psoriasis, atopic dermatitis, actinic keratosis, actinic pass or be used to treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal detachment, and for treating and preventing scarring such as keloid, adhesion and hypertrophic or hypertrophic burn and side in the second or and seed or the sequences used in the
                                                                                                                                                                                         Human; ribozyme therapy; hairpin ribozyme; hammerhead ribozyme; recognition site; target; ribozyme binding site; eye disease; vulnerary; proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin; MWP; matrix metalloproteinase; growth factor; reductase; scarting; cytostatic; antipsoriatic; dermatclogical; antiesborrheic; antidiabetic; virucide; antipsickling; ophthalmological; keratolytic; gene therapy; viral wart; basal cell carcinoma; seborrheic wart; vitreoretinopathy; scar; sickle cell retinopathy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent kinases.
                                                                                                                                                 Cell-cycle dependent kinase cdk8 ribozyme binding site SEQ ID NO:956.
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AAH58532
ID AAH58532 standard; DNA; 19 BP.
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                                                                                                          10-SEP-2001 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
Synthetic.
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                                                              AAH58532;
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Assessing susceptibility to syndromes including migraine with aura, depression and anxiety - by detecting variant alleles in genes for dopaminergic receptors or transporter, also treatment using agents that antagonise binding of dopamine to these proteins.

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The present sequence is a primer for the dopaminergic gene DRD5, which can be used in a novel method assess susceptibility to a syndrome having symptoms of migraine with aura (MMA), depression and/or anxiety. The method comprises detecting a variant allele of at least 1 dopaminergic gene. Analysis of variant dopaminergic gene alleles may also differentiate between patients with MMA, and those with migraine without aura (MO) or other conditions, e.g. stroke, that produce similar symptoms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mouse C/EBP beta phosphorothioate antisense oligonucleotide, SEQ ID:127.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mouse; murine; C/EBP beta; CCAAT/enhancer-binding protein beta; C/EPB2; LAP, TCF5; CRP2; NFIL6; IL6DBP; NF-M; AGP/EBP; Apc/EBP; transcription factor; tissue development; cellular function; proliferation; differentiation; hormone responsiveness;
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Best Local Similarity 93.8°
Matches 15; Conservative
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Gaps

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3.4%; Score 14.4; DB 1; Length 19; 93.8%; Pred. No. 2e+02; ive 0; Mismatches 1; Indels

GGTGAAGGACCTGAGC 308

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Best Local Similarity 93.8 Matches 15; Conservative

Query Match

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/mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides. All 2' MOE
cytosines are 5-methylcytosine"
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/mod_base= CTHER
/mod= "2'-methoxyethyl (2'-MOE) nucleotides. All 2' MOE
cytosines are 5-methylcytosine"
                                                                                                                                                                                                                                                                                                                                                                    Novel antisense compound targeted to nucleic acids encoding human or mouse CCAAT/enhancer binding protein (C/EBP) beta, useful in vitro for inhibiting expression of human or mouse C/EBP beta in cells/tissues.
oxidative stress response; IL-6 signalling mediator; interleukin-6; carbohydrate metabolism; immunity; Thi response; female fertility; glucomeogenesis; ovarian; cancer; tumour formation; type II; diabetes; infection; inflammation; expression inhibition; phosphorothioate; antisense oligonucleotide; ss.
                                                                                                     /mod_base= OTHER
/note= "Phosphorothioate linkages"
1. 5
/*tag= b
                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                          14-JUN-2000; 2000US-00593711.
                                                                                                                                                                                                                                                                                            14-JUN-2000; 2000US-00593711.
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                                                                                                                                                                           16. .20
                                                                                        1. .20
/*tag=
                                                                                                                                                                                                                                                                                                                                  BP, Butler MM,
                                                                                                                                                                                                                                                                                                               (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-214451/27.
                                                                                Key
modified_base
                                                                                                                                                                             modified_base
                                                                                                                              modified base
                                                              Mus musculus
                                                                                                                                                                                                                                     US6271030-B1
                                                                                                                                                                                                                                                       07-AUG-2001.
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Sequences ABL94252-ABL94476 represent antisense oligonucleotides targeted to the human or mouse CCAAT/enhancer-binding protein alpha (C/BBP alpha) and which inhibit its expression. The antisense oligonucleotides were designed to target different regions of the human and/or mouse C/BBP alpha RNA, and were analysed for their effect on C/BBP alpha mENA levels alpha RNA, and were analysed for their effect on C/BBP alpha mENA levels by quantitative real-time PCR. The C/BBP family of proteins are a family of transcription factors which regulate the expression of a wide range of commercial insue development, cellular function, cellular control normal lissue development, cellular function, cellular correspondents of proliferation and functional differentiation. C/BBP beta (also known as C/BPBP, LAP, TCF5, CRP2, NRILG, ILGBP, NP, AGP/BBP and Apc/BBP) compared to primarily regulates hormone responsiveness and oxidative stress responses compared to be involved in carbohydrate metabolism, immunity, the Thi cresponse, female fertility and gluconeogenic pathways. C/BBP beta is thought to be involved in carbohydrate metabolism, immunity, the Thi cresponse, female fertility and gluconeogenic pathways. C/BBP beta is capressed in the liver, lung, spleen, kidney, brain, and testils, with the highest expression found in the lung. It is also expressed at a higher compared level is of glucose, indicating that it is involved in the level sof glucose, indicating that it is involved in the celevated levels of glucose, indicating that it is involved in the conditions associated with C/BBP beta expression, gunt as cancer (particularly ovarian cancer), tumour formation, diabetes (particularly veryian cancer), infection, or inflammation Claim 1; Col 47-48; 69pp; English

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3.4%; Score 14.4; DB 1; Length 20; 93.8%; Pred. No. 2.2e+02;
Sequence 20 BP; 1 A; 6 C; 8 G; 5 T; 0 U; 0 Other;
                                                      Query Match
Best Local Similarity
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Sequences ABL94252-ABL94476 represent antisense oligonucleotides targeted to the human or mouse CCAAT/enhancer-binding protein alpha (C/BBP alpha) gene, which inhibit its expression. The antisense oligonucleotides were designed to target different regions of the human and/or mouse C/BBP alpha RNA, and were analysed for their effect on C/BBP alpha mRNA levels by quantitative real-time PCR. The C/BBP family of proteins are a family of transcription factors which regulate the expression of a wide range of genes that control normal tissue development, cellular function, cellular /*tag= c/mod_base OTHER /mod_base OTHER /note= "2'-methoxyethyl (2'-MOE) nucleotides. All 2' MOE cytosines are 5-methylcytosine" Mouse C/EBP beta phosphorothioate antisense oligonucleotide, SEQ ID:128. Mouse; murine; C/EBP beta; CCAAT/enhancer-binding protein beta; C/EPB2; LAP, TCF5; CRP2; NFIL6; IL6DBP; NF-M; AGP/EBP; Apc/EBP; transcription factor; tissue development; cellular function; proliferation; differentiation; hormone responsiveness; oxidative stress respons; IL-6 signalling mediator; interleukin-6; carbohydrate metabolism; immunity; Thi response; female fertility; gluconeogenesis; ovarian; cancer; tumour formation; type II; dlabetes; infection; inflammation; esp. /note= "2'-methoxyethyl (2'-MOE) nucleotides. All 2' cytosines are 5-methylcytosine" 16. .20 Novel antisense compound targeted to nucleic acids encoding human or mouse CCAAT/enhancer binding protein (C/EBP) beta, useful in vitro finhibiting expression of human or mouse C/EBP beta in cells/tissues. Gaps ö Indels /mod_base= OTHER /note= "Phosphorothicate linkages" ï Mismatches Location/Qualifiers 1. 20 /*tag= a Claim 1; Col 49-50; 69pp; English. /*tag= b /mod_base= OTHER Monia BP, Butler MM, Wyatt J; ö 14-JUN-2000; 2000US-00593711. 383 CGACGACGGCGCCAAG 398 ABL94362 standard; DNA; 20 29-JUL-2002 (first entry) 16 CGACTACGCCGCCAAG Matches 15; Conservative (ISIS-) ISIS PHARM INC. WPI; 2002-214451/27. Key modified_base modified_base modified_base 14-JUN-2000; Mus musculus 07-AUG-2001. ABL94362; ABLO94362/CABLO9 ABLO94362/CABLO9 ABLO94362/CABLO9 ABLO94362/CABLO9 ABLO96/CABLO9 ABLO96/CABLO9 ABLO96/CABLO9 ABLO96/CABLO9 ABLO96/CABLO9 ABLO96/CABLO9 ABLO96/CABLO9 ABLO96/CABLO9 ABLO96/CAB 쉱 ઠે

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proliferation and functional differentiation. C/EBP beta (also known as C/EBP) LAP, TCFS, (REP2, NETL6, IL6DBP, NR-M, AGP/EBP) and App/EBP)

primarily regulates hormone responsiveness and oxidative stress responses and is a mediator of IL-6 (interlukin-6) signalling. C/EBP beta is thought to be involved in carbohydrate metabolism, immunity, the Thi response, female fertility and gluconeoganic pathways. C/EBP beta is expressed in the liver, lung, spleen, kidney, brain, and testis, with the lighest expression found in the lung. It is also expressed at a higher level in malignant ovarian tissue compared with normal ovarian tissue, and its expression in pancreas is upregulated in response to chronically and its expression in pancreas is upregulated in response to chronically impairment of insulin secretion in type II diabetes. The oligonucleotides of the invention are useful for diagnosis, prevention and treatment of confittions associated with C/EBP beta expression, such as cancer confittions associated with C/EBP beta expression, such as cancer (particularly ovarian cancer), tumour formation, diabetes (particularly type II diabetes), infection, or inflammation

Sequence 20 BP; 1 A; 6 C; 8 G; 5 T; 0 U; 0 Other;

ö 3.4%; Score 14.4; DB 1; Length 20; 93.8%; Pred. No. 2.2e+02; ive 0; Mismatches 1; Indels 383 CGACGACGCGCCAAG 398 ហ Local Similarity 93.8 nes 15; Conservative 20 Query Match Best Loca Matches ઠે g

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RESULT 105 AAV32912/c ID AAV32912 standard; DNA; 21

AAV32912;

(first entry) 26-OCT-1998

Bovine lactoferrin cDNA primer 1.

PCR; primer; amplification; pepsin; gastrointestinal tract; milk; Aspergillus niger beta-galactosidase gene; lactase intolerance; cheese making; chymosin; bovine lactoferrin cDNA; ss.

Synthetic

WO9829536-A2

09-JUL-1998

97WO-IB001658 29-DEC-1997; 96US-00775842. 31-DEC-1996; (NEXI-) NEXIA BIOTECHNOLOGIES INC.

Kabel JJ, Amantea GF Turner JD, Eino M, Karatzas CN, 

WPI; 1998-388118/33.

- can be useful Synthetic beta-galactosidase inactive in milk but active in vivo chemically activated and used to treat lactose intolexance, also in cheese production.

Example 1; Page 13; 38pp; English.

Primers 1 and 2 (AAV32913) were used in a PCR reaction to amplify the bovine lactoferrin cDNA. The PCR product was used as a tail which was fused through a pepsin recognition site to the 3' end of the Aspergillus niger beta-galactosidase gene. The invention provides a synthetic beta-galactosidase which differs from the natural occurring enzyme in being inactive in milk but capable of being activated by a chemical or condition naturally present in the gasrointestinal tract of humans. The design of this synthetic enzyme comprises of a tail domain fused to the

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beta-galactosidase through a cleavage site. The presence of the tail domain renders the enzyme inactive and it can also be used as a purification handle. The synthetic beta-galactosidase is claimed to be able to hydrolyse lactose in vivo to overcome lactase intolerance and thereby reduce associated gastrointestinal disorders. The synthetic betaglactosidase is also claimed to be useful in cheese making whereby it is activated by chymosin when added to milk
                                                                                                                                                                                                                                                                                                                                                Human; variant thrombospondin 1; variant thrombospondin 4; SNP; polymorphism; vascular disease; coronary artery disease; forensics; myocardial infarction; atherosclerosis; stroke; venous thromboembolism; pulmonary embolism; paternity test; ds.
                                                                                                                                            ;
                                                                                                                   Query Match 3.4%; Score 14.4; DB 1; Length 21; Best Local Similarity 93.8%; Pred. No. 2.5e+02; Matches 15; Conservative 0; Mismatches 1; Indels
                                                                                            Sequence 21 BP; 4 A; 5 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                          Human gene single nucleotide polymorphism #16.
                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
replace(11,G)
/*tag= a
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                                                                                                                                                                   192 ATCCACTGCTCGGTGA 207
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                                                                                                                                                                                                                                                     AAF95255 standard; DNA; 21
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Variation
                                                                                                                                                                                                                                                                            AAF95255;
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Mccarthy JJ /*tag= a /standard_name= "single nucleotide polymorphism" Gargill M, Ireland JS, Bolk S, Daley GQ, (WHED ) WHITEHEAD INST BIOMEDICAL RES. (MILL-) MILLENNIUM PHARM INC. 10-SEP-1999; 99US-0153357P. 26-JUL-2000; 2000US-0220947P. 16-AUG-2000; 2000US-0225724P. 07-SEP-2000; 2000WO-US024503 WO200118250-A2 15-MAR-2001. Lander ES,

Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and WPI; 2001-226749/23

Example; Page 48; 242pp; English.

atherosclerosis.

The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various obly morphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thromboembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also

Sequence 21 BP; 6 A; 6 C; 8 G; 1 T; 0 U; 0 Other;

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The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various bolymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, cornary heart disease, stroke, peripheral vascular diseases, venous thromboembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; variant thrombospondin 1; variant thrombospondin 4; SNP; polymorphism; vascular disease; coronary artery disease; forensics; myocardial infarction; atherosclerosis; stroke; venous thromboembolism; pulmonary embolism; paternity test; ds.
useful in forensics, paternity testing, genetic analysis and phenotype correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gargill M, Ireland JS, Bolk S, Daley GQ, Mccarthy JJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and
                                                                                                                                             Gaps
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/standard_name= "single nucleotide polymorphism"
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                                                                                                    3.4%; Score 14.4; DB 1; Length 21; 93.8%; Pred. No. 2.5e+02; tive 0; Mismatches 1; Indels
                                                                     Sequence 21 BP; 7 A; 5 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                     Human gene single nucleotide polymorphism #1169,
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                                                                                                                                                                                204 GIGAAAGCAGAGACT 219
                                                                                                                                                                                                                                                                                                          AAF96408 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-SEP-1999; 99US-0153357P.
26-JUL-2000; 2000US-0220947P.
16-AUG-2000; 2000US-0225724P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-SEP-2000; 2000WO-US024503
                                                                                                                                                                                                                   GTGAATGCAGAGAACT 21
                                                                                                                                                                                                                                                                                                                                                                                    06-JUN-2001 (first entry)
                                                                                                                          Best Local Similarity 93.8
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Key
Variation
                                                                                                                                                                                                                                                                                                                                               AAF96408;
                                                                                                          Query Match
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                                 Gaps
                                                                                                                                                                                                                                                                                            yak milk; alpha-lactoalbumin; beta-lactoglobulin; alpha S1-casein; alpha S2-casein; beta-casein; kappa-casein; lactoferritin; ss.
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Score 14.4; DB 1; Length 21; Pred. No. 2.5e+02;
                               1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 6 A; 4 C; 11 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 5 (disclosure); 41pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Seven kinds of yak milk protein gene sequence.
   3.4%; Score 14.4; D
Local Similarity 93.8%; Pred. No. 2.5e
les 15; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                 Yak milk protein gene related oligo, F30.
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                                                                301 ACCTGAGCCCCGGGGA 316
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28 AGGGCTGGGACGAAGA 43
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                                                                                            1 Accreaeccceaeca 16
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                                                                                                                                                                        ADE64663 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-OCT-1996 (first entry)
                                                                                                                                                                                                                                   (first entry)
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Matches 15, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-741796/81.
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      Query Match
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AC AAT13
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DT 29-00
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The present sequence is a PCR primer for plasmid pBlue-TH6, which was used in the construction of the expression plasmid pEdHCkappa-TH6. pEdHCkappa-TH6 was prepd. by inserting a gene encoding the light-chain [LC) variable region of human anti-TSH antibody (Ab) into pEdHCkappa, an expression vector for a Ab LC. pEdHCkappa-TH6, an expression vector for the prodn. of an Ab LC in an animal host cell, contains 5'3' a SV40 promoter, and base sequences encoding dihydrofolate reductase, Ab LC signal sequence and Ab LC variable and constant regions. pEdHCkappa-TH6 along with the equivalent heavy chain expression vector pEdHCG1-TH8 can be used for the prepn. of an Ab mol. in an animal host cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Expression vectors for antibody (Ab) heavy and light chains - introduced concurrently into animal host cell to produce Ab mol. which is secreted
                                                                      Plasmid; pBlue-TH6; construction; expression plasmid; primer; pBdHKappa-TH6; light chain; variable region; human; antibody; TSH; thyroid stimulating hormone; animal host cell; SV40 promoter; dihydrofolate reductase; PCR; polymerase chain reaction; ss.
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Plasmid pBlue-TH6 PCR primer TSEVK1FOR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (TOYJ ) TOSOH CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      concurrently in in supernatant.
                                                                                                                                                                                                                                                                                                                                                                           JP08051995-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-AUG-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-AUG-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               27-FEB-1996.
                                                                                                                                                                                                                                                                                           Synthetic.
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Crasscripts expressed in Xenomice. The products were used for repertoire transcripts expressed in Xenomice. The products were used for repertoire analysis. The specification describes a transgenic non-thuman mammal which has genome modifications that comprise an inactivated endogenous Ig locus, so that the mammal does not display normal B-cell development. The conditions also has an inserted human heavy chain Ig locus in germline configuration, the human heavy chain Ig locus in micro constant region and regulatory and switch sequences, human J-H genes and an inserted human wappa constant region and regulatory and switch sequences, human J-H genes, and human W-H genes and an inserted human wappa continuing to Ight chain Ig locus in germline configuration, the human kappa light chain Ig locus comprising a human Nappa constant region, J-kappa genes, where the number of V-H and V-kappa genes inserted are selected to restore normal B-cell development in the mammal. The cransgenic animals have a near complete human Ig locus, including both a transgenic animals have a near complete human Iglocus, including both a cused for the production of human antibodies when exposed to particular antigens e.g. when exposed to human antibodies when respectively

New transgenic non-human mammals - having an inactivated immunoglobulin locus and a near complete human immunoglobulin locus, used for production

Disclosure, Page 30, 128pp, English.

of human antibodies

Green L;

Mendez M,

Klapholz S,

Jakobovits A, Kucherlapati R,

(ABGE-) ABGENIX INC.

WPI; 1998-33314/29.

97WO-US023091 96US-00759620

03-DEC-1997; 03-DEC-1996; ö

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98; PCR; primer; amplification; human; epidermal growth factor receptor;
tumour; BGR; transforming growth factor alpha; TGF-alpha.
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                                                                                                                                                                                                                                                                                                               Score 14.4; DB 1; Length 22; Pred. No. 2.7e+02;
                                                                                                                                                                                                                                                                                                                               1; Indels
                                                                                                                                                                                                                                                                                               Sequence 22 BP; 4 A; 5 C; 10 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Jia X;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gallo M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human universal VH primer MG-30.
                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                            AAV68617 standard; DNA; 22 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98WO-US009160.
                                                                                                                                                                                                                                                                                                                                                  263 GGTGCACCTGGAGCAG 278
                                                                                                                                                                                                                                                                                                                3.4%;
                                                                                                                                                                                                                                                                                                                                                              GTGCAGCTGGAGCAG 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   97US-00851362
                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 93.89
Marches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Yang X,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ABGE-) ABGENIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Jakobovits A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9850433-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                             30-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-NOV-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                              AAV68617;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; immunoglobulin; Ig; transgenic; non-human mammal;
inactivated endogenous Ig locus; B-cell development;
human heavy chain Ig locus; micro constant region; J-H; D-H; V-H gene;
kappa light chain Ig locus; kappa constant region; J-kappa gene; V-kappa;
production; antibody; PCR primer; 88.
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Universal human VH PCR primer MG-30.

Homo sapiens WO9824893-A2

Synthetic

11-JUN-1998

23-SEP-1998 (first entry)

AAV42250;

Gaps

; 0

3.4%; Score 14.4; DB 1; Length 22; 93.8%; Pred. No. 2.7e+02; tive 0; Mismatches 1; Indels

297

GGCACCAAGCTGGTGA

282

ò 셤

Local Similarity 93.8 les 15; Conservative

Query Match Matches 22 GGCACCAAGCTGGAGA 7

BP

AAV42250 standard; DNA; 22

AAV42250

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                                                                                                                                                   The primers AAV68617-V68618 were used to produce anti-epidermal growth factor receptor (EGF-r)-antibodies. The antibodies can be administered therapeutically to patients (human or veterinary) to treat solid tumours EGF-r is overexpressed on many human solid tumour types, and the fully human antibodies (i.e. comprising (i) and (ii)) inhibit both epidermal growth factor (EGF) and transforming growth factor alpha (TGF-alpha) binding to EGF-r (Known to lead to cellular proliferation and tumour growth). They can prevent tumour cell growth and, in combination with an antineoplastic agent, may eradicate established tumours. The fully human antibodies can minimise the immunogenic and allergic responses intrinsic to previous mouse/rat or mouse/rat-derived antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sox-9; bone regeneration; cartilage regeneration; campomelic dysplasia; gene therapy; sex reversal; primer; single strand conformation polymorphism; SSCP; PCR; polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SSCP primers 534 (AAT30310), 661 (AAT30311), 687 (AAT30312), 854 (AAT30313), 836 (AAT30314) and 1018 (AAT30315) were used for SSCP analysis of the SOX-9 gene (see also AAT30309) in campomelic dysplasia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New isolated SOX-9 genes - used to develop prods. for the promotion or suppression of bone or cartilage differentiation of growth.
                                 Humanised antibodies against epidermal growth factor receptor, EGF-r useful to treat solid tumours whilst inducing reduced immunogenic or allergic effects compared to mouse or mouse-derived antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gape
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.4%; Score 14.4; DB 1; Length 22;
93.8%; Pred. No. 2.7e+02;
tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 22 BP; 4 A; 5 C; 10 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 42; 64pp; English.
                                                                                                                      Example 3; Page 96; 143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                263 GGTGCACCTGGAGCAG 278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-NOV-1994; 94AU-00009714.
05-DEC-1994; 94AU-00009835.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GGTGCAGCTGGAGCAG 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAT30310 standard; cDNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Goodfellow PN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ouery Match
Best Local Similarity 93.8%
Matches 15, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOX-9 SSCP primer 534.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1996-277777/28.
WPI; 1999-034712/03.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-AUG-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAT30310;
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AAT3031
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                                                                                                                                                               ö
(CD) patients. Primers were designed to amplify the known coding sequence and intron/exon junctions. Unique SSCP conformers were cloned and sequenced. Alterations in SOX-9 can cause both CD and male-to-female sex reversal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human glial cell derived neurotrophic factor and its derivatives and use.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                        derived neurotrophic factor; GDNF; PCR; primer; 88;
                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                         Human glial cell derived neurotrophic factor (GDNF) PCR primer #44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
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3.3%; Score 14.2; DB 1; Length 19;
Best Local Similarity 84.2%; Pred. No. 2.2e+02;
Matches 16; Conservative 0; Mismatches 3; Indel8
                                                                                                                                Length 19;
                                                                                                                                                               3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19 BP; 3 A; 8 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                               Sequence 19 BP; 7 A; 2 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                              Query Match
3.3%; Score 14.2; DB 1;
Best Local Similarity 94.2%; Pred. No. 2.2e+02;
Matches 16; Conservative 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (YISH-) YISHENG BIOLOGICAL PHARM CO LTD SHUHAI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 6; Page 4 (Claims); 28pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       228 GCCAAATCGGGAGGCTGCT 246
                                                                                                                                                                                                350 GCTCTACAGCGACTTCCTC 368
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19 GCGGAATCGGCAGGCTGCT 1
                                                                                                                                                                                                                      19 GTTCTTCACCGACTTCCTC 1
                                                                                                                                                                                                                                                                                                                 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-JAN-2001; 2001CN-00107450.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11-JAN-2001; 2001CN-00107450.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAQ73805 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                 ACA96850 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Zhou S, Zheng Z, Feng H;
                                                                                                                                                                                                                                                                                                                                                                               24-JUL-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nervous system disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-000523/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                             Human, glial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CN1364812-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-AUG-2002.
                                                                                                                                                                                                                                                                                                                                                   ACA96850;
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AAQ73805/c
ID AAQ7380
XX
                                                                                                                                                                                                                                                                                                   ACA96850,
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                        8X33333
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(revised) (first entry)

25-MAR-2003 22-MAY-1995

AAQ73805;

Aspergillus aculeatus; ss

Pectin lyase;

Aspergillus aculeatus.

WO9421786-A1

29-SEP-1994

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which are complementary to at least a portion of the human osteopontin (OPN) cDNA sequence (AAX29191). The antisense sequences are used to prevent restenosis in tissue, particularly coronary arterial tissue, especially where the patient is undergoing angioplasty, particularly percutaneous trans-luminal coronary angioplasty or directional coronary atherectomy. They prevent secretion of osteopontin by monocytes and macrophages which infiltrate to sites of inflammation following surgery osteopontin probably causes restenosis by inducing coronary artery smooth muscle cells (CASMC) to migrate to, and proliferate at, angioplasty injury sites. Sequences AAX29177-178 represent RT-PCR primers specific
                                                                                                                                                                                                                                                                                                              relates to antisense osteopontin oligonucleotide sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human, mitochondrial phosphoenolpyruvate carboxykinase; PBPCK-M; PCK2; PEPCK-mitochondrial; mtPEPCK, antisense oligonucleotide; modulation; phosphorothioate; inhibition; diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human mtPEPCK phosphorothioate antisense oligonucleotide SEQ ID NO:27
                                                                                                                                                                                                                New osteopontin antisense sequences - useful to treat restenosis, particularly following vascular surgery.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /*tag= a
/note= "phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 5 A; 6 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 14.2; DB 1;
Pred. No. 2.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                         & HUMAN SERVICES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
1. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            for human osteopontin cDNA sequence
                                                                                                                                               Panda DK;
                                                                                                                                                                                                                                                                            Example 1; Page 28; 72pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 CACCAGTCTGATGAGTCTC 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          48 CACCACTCAGAGGAGTCTC 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3.3%;
                                 98WO-US016569
                                                                     97US-0054967P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16; Conservative
                                                                                                                                               Kundu GC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ISIS-) ISIS PHARM INC.
                                                                                                         (USSH ) US DEPT HEALTH
                                                                                                                                                                                  WPI; 1999-190049/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
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modified_base
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Synthetic.
                                                                                                                                                 Mukherjee AB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31-MAY-2000
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                                                                       07-AUG-1997;
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                                 07-AUG-1998;
18-FEB-1999.
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AAZ95339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAQ73789-Q73822 are partial DNA sequences, one or more of which can be used to encode enzymes having pectin lysae (PL) activity. The Aspergillus sequences by and the coresponding DNA sequence, from which these partial sequences were derived are shown in AAK60881 and AAQ73823 respectively. These PL enzymes degrade plant cell wall components, and can therefore be used to reduce the viscosity of fruit juices. They can also be used for the production of antibodies. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Osteopontin; antisense; restenosis; coronary arterial tissue; CASMC; inflammation; coronary artery smooth muscle cell; angioplasty; human; OPN; RT-PCR; primer; ss.
                                                                                                                               cell wall degradation; reducing fruit juice viscosity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New pectin lyase enzyme from Aspergillus aculeatus - used for the degradation of plant cell wall components, esp. for reducing the viscosity of fruit juices.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Christgau S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
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84.2%; Pred. No. 2.4e+02;
ive 0; Mismatches 3; Indels
                                                                                           Aspergillus aculeatus pectin lyase partial DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human osteopontin (OPN) specific RT-PCR primer hOPN-L
                                                                                                                                                                                                                                                                                                                                                                                                                                   Kofod LV, Kauppinen MS, Andersen LN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 3 A; 8 C; 6 G; 3 T; 0 U; 0 Other;
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Claim 2; Page 47; 65pp; English.

93DK-00000279. 94WO-DK000105

12-MAR-1993; 28-OCT-1993;

11-MAR-1994;

(NOVO ) NOVO-NORDISK AS

WPI; 1994-317007/39.

Dalboge H, Kofo Heldt-Hansen HP;

316 ACCGCGTGCTGGCGGCGA 334

Conservative

Local Similarity

Query Match

16;

Best Loca Matches

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Accaccificatectecce

20

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AAX29178 standard; DNA; 20

RESULT 115

AAX2917

(first entry)

18-JUN-1999

AAX29178;

Synthetic. Homo sapiens. WO9907844-A2

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Gaps

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New antisense compound targeted to a nucleic acid molecule encoding human mitochondrial phosphoenolpyruvate carboxykinase useful for treating a human with a mitochondrial phosphoenolpyruvate carboxykinase-associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAZ95320 to AAZ95359 represent antisense oligonucleotides targeted to a nucleic acid molecule encoding human mitochondrial phosphoenolpyruvate carboxykinase (also known as PEPCK-mitochondrial, PEPCK-M; PCK2 and mtPEPCK), where the oligonucleotide specifically hybridise with and inhibit the expression of human mtPEPCK. The antisense oligonucleotides can be used for inhibiting the expression of mtPEPCK in human cells or tissues in vitro and can also be used for treating an animal, particularly a human suspected of having or being prone to a condition odisease associated with expression of mtPEPCK. They can also be used in diagnostics and as research reagents in sandwich and other assays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 3 A; 7 C; 7 G; 3 T; 0 U; 0 Other;
Butler MM
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 3; Col 39; 32pp; English.
Cowsert LM,
                                                                                                WPI; 2000-205209/18.
Mckay R,
                                                                                                                                                                                                                                                                                                                                                       disease.
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Ouery Match 3.3%; Score 14.2; DB 1; Length 20; Best Local Similarity 84.2%; Pred. No. 2.4e+02; Matches 16; Conservative 0; Mismatches 3; Indels
                                                                                                                        136 CCCGCCTGGCGGTGGAGGC 154
                                                                                                                                                                      2 ccadccrddcadrdcaddc 20
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Gaps .; 0

> Human B7-1 antisense oligonucleotide SEQ ID NO: 154. AAF32957 standard; DNA; 20 (first entry) 23-MAR-2001 AAF32957;

BP.

Human, mouse; B7-1; B7-2; antisense; PCR primer; inflammation; autoimmune disorder; phosphorothloate backbone; ss. WO200074687-A1. Homo sapiens. 14-DEC-2000. 

25-MAY-2000; 2000WO-US014471. 99US-00326186 (ISIS-) ISIS PHARM INC. 04-JUN-1999;

Bennett CF, Vickers TA,

Karras JG

WPI; 2001-049991/06.

Novel compound for diagnosing, preventing and treating immune disorders, comprising an oligonucleotide that specifically hybridizes with a nucleic acid sequence encoding B7 protein.

Example 12; Page 76; 162pp; English.

The present invention provides sequences of antisense oligonucleotides targeted at the murine and human B7-1 and B7-2 coding and mRNA sequences. The antisense sequences have phosphorothicate backbones and some nucleotides are 2'-methoxyethoxy residues. The sequences can be used in the treatment of inflammatory and autoimmune disorders, including asthma,

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juvenile diabetes mellitus, myasthenia gravis, Graves' disease, rheumatooid arthritis, allograft rejection, inflammatory bowel disease, multiple sclerosis, psoriasis, systemic lupus erythematosus, contact dermatitis, rhinitis, allergies and cancer
                                                                                                                                                                  Gaps
                                                                                                                                                                .
                                                                                                                             Length 20;
                                                                                                                                                            3; Indels
                                                                                        Sequence 20 BP; 6 A; 8 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                         3.3%; Score 14.2; DB 1;
84.2%; Pred. No. 2.4e+02;
tive 0; Mismatches 3;
                                                                                                                                                                                                   398 GAAGGICTICTACGTGATC 416
                                                                                                                                                                                                                                      19 gaaggrerrcrrcgraagc 1
                                                                                                                     Query Match
Best Local Similarity 84.29
Matches 16; Conservative
   8888888
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AAC84282 standard; DNA; 20

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(first entry)

19-MAR-2001

AAC84282;

New signal transduction nucleic acids and encoded proteins useful for regulating phytohormone expression, including ethylene, auxins, cytokinins and gibberellin, to provide control of plant response to environmental stresses. Zea mays; maize; signal transduction protein; phytohormone; ethylene; auxin; cytokinin; gibberellin; immunogen; PCR primer; ss. Signal transduction cDNA amplifying primer. (PION-) PIONEER HI-BRED INT INC. 99US-0134292P. 99US-0142996P. 28-APR-2000; 2000WO-US011687. WPI; 2001-031929/04. WO200070059-A2. Helentjaris TG; 14-MAY-1999; 08-JUL-1999; 23-NOV-2000. Zea mays. 

The invention provides Zea mays signal transduction proteins and encoding nucleotide sequences. The nucleic acids are useful for regulating explanating captences. The nucleic acids are useful for regulating captences. The mack awxins, cytokinins, and gibberslin, to effect developmental changes in plants and provide control of plant response to environmental stresses. They may also be used as probes or amplification primers in the detection, quantitation or isolation of gene transcripts, for detecting mutations in the gene, for monitoring upregulation of expression or changes in enzyme activity in cremening assays of compounds, for detection of any number of allelic variants, or for site-directed mutagenesis in eukaryotic cells. They may further be used for recombinant expression of their encoded polypeptides, as immunogens in the preparation or screening of antibodies, and in sense or antisense suppression of genes in a host cell, tissue or plant. The proteins may be used in assays for enzyme agonists or antisquists, as immunogens or antigens to obtain antibodies specifically immunoreactive with the proteins. The present sequence represents a PCR primer used for amplifying the cDNA encoding a signal transduction protein

Example; Page 111; 126pp; English.

Seguence 20 BP; 3 A; 8 C; 5 G; 4 T; 0 U; 0 Other;

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modified_base
                                                                                                                                                                                                                                                                                                                                                                                      modified_base
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                                                                                                                                                                                                                                                     30-OCT-2002
                                                                                                                                                                                                                                                                                              Homo sapiens
                   Cowsert LM;
                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                          AAD40675;
                                                                                                                                                                                                                 RESULT 120
AAD40675/c
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             Gaps
                                                                                                             Human; hepsin; antisense compound; antisense therapy; antisense; phosphorothioate backbone; ss.
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Query Match 3.3%; Score 14.2; DB 1; Length 20; Best Local Similarity 84.2%; Pred. No. 2.4e+02; Matches 16; Conservative 0; Mismatches 3; Indels
                                                                                                   Human hepsin antisense oligonucleotide, ISIS 107131.
                                                                                                                                                                                                                                                                                                                                      /rtag= k
/mod_base= m5c
16. .20
/*tag= c
/mod_base= OTHER
/mod_base= OTHER
                                                                                                                                                                                *tag= b
"tagbas= OTHER
"note= "2'methoxyethyl nucleotides"
                                                                                                                                                         . . . 20
*tag= a
mod base= CTHER
note= "Phosphorothioate backbone"
                                                                                                                                                  ocation/Qualifiers
                       139 GCCTGGCGGTGGAGGCCGG 157
                                                                                                                                                                                                           /*tag= d
/mod_base= m5c
                               20 GCCTGGCGGTGGAAACCTG 2
                                                                                                                                                                                                                                                                                  *tag= h
mod_base= m5c
                                                                                                                                                                                                                                                                                                  *tag= i
mod_base= m5c
                                                                                                                                                                                                                                                                                                                     *tag= j
mod_base= m5c
                                                                                                                                                                                                                                               *tag= f
mod_base= m5c
                                                                                                                                                                                                                                                                 *tag= g
mod_base= m5c
                                                                                                                                                                                                                              '*tag≈ e
mod_base= m5c
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                                                           AAD40857/c
ID AAD40857 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                         14-DEC-2001; 2001WO-US048341.
                                                                                                                                                                                                                                                                                                                                                                                                                                    20-DEC-2000; 2000US-00742482.
                                                                                       30-OCT-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                *tag= 1
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                                                                                                                                                   Key
modified_base
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                                                                                                                                 Homo sapiens.
Synthetic.
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                                                                            AAD40857;
                                                     RESULT 119
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                                                                                                                                                                                                                                                                                                                                             The invention relates to antisense compounds, compositions and methods for modulating the expression of hepsin. The compositions comprise antisense compounds, particularly antisense oligonucleotides, targetted to nucleic acids encoding hepsin. The antisense compound is useful for inhibiting the expression of hepsin in human cells or tissues. It is also useful for treating an animal having a disease or condition associated with hepsin, by inhibiting expression of hepsin. It is useful for diagnostics, therapeutics, prophylaxis and as research reagents and kits. It is also used in antisense therapy. The present sequence is an entisense oligonuclectide targetted to human hepsin DNA. This sequence is used in the exemplification of the invention
                                                                                                                                                                      Novel antisense compound targeted to nucleic acids encoding human hepsin, useful for inhibiting the expression of hepsin in human cells or tissues, and for treating humans having a disease associated with human hepsin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; antisense; hepsin; inflammation; tumour; gene therapy; cytostatic; phosphorothioate backbone; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
3.3%; Score 14.2; DB 1; Length 20;
Best Local Similarity 84.2%; Pred. No. 2.4e+02;
Matches 16; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human hepsin antisense oligonucleotide, ISIS 107131.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mod_base= OTHER
/note= "2'methoxyethyl nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 4 A; 9 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                       Claim 3; Page 97; 100pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          308 CCCCGGGGACCGCGTGCTG 326
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20 CTCCGGGGACTGGGTG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /*tag= e
/mod_base= m5c
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/mod_base= m5c
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/mod_base= m5c
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/mod_base= m5c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ...5
*tag= b
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
(ISIS-) ISIS PHARM INC
                                                                                                                 WPI; 2002-519882/55.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Key
modified_base
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Human RIP2 antisense oligonucleotide ISIS #104251
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Human; receptor interacting protein; RIP2; antisense; gene therapy; phosphorothicate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                              01-AUG-2001; 2001US-00920663
                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-AUG-2001; 2001US-00920663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ward DT, Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-673017/72.
                                                                                        Key
modified_base
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                                                                                                                                                                                                                                                                                                                                                                                       US6426221-B1
                                                         Homo sapiens.
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                           30-JUL-2002.
The invention relates to an antisense compound 8-30 nucleobases in length targetted to a nucleic acid molecule encoding human hepsin. The antisense compound inhibits the expression of human hepsin. The antisense compound or the pharmaceutical composition of useful for treating animals and human having a disease or condition associated with the expression of hepsin, e.g. inflammation or tumour growth. The antisense compounds are useful also for diagnositis, prophylaxis (e.g. to prevent or delay infection, inflammation or tumour formation) or as research reagents and kits. The method is useful for modulating, specifically inhibiting the expression of hepsin which may be used in research, e.g to distinguish between functions of various members of a biological pathway. The invention is used in gene therapy. The present sequence is human hepsin antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                        New antisense oligonucleotides that modulate (particularly inhibit) human hepsin, useful for treating a disease or condition associated with the expression of hepsin, e.g. inflammation or tumor growth.
                                                                                              *tag= k
/mod_base= m5c
16, .20
/*tag= c
/mod_base= OTHER
/note= "2'methoxyethyl nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 4 A; 9 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                   Example 15; Page 82; 101pp; English.
                                     *tag= i
mod_base= m5c
                                                                    *tag= j
mod_base= m5c
         *tag= h
mod_base= m5c
                                                                                                                                                                       /*tag= l
/mod_base= m5c
                                                                                                                                                                                                                                              14-DEC-2001; 2001WO-US048431.
                                                                                                                                                                                                                                                                 20-DEC-2000; 2000US-00742703
                                                                                                                                                                                                                                                                                                                     Cowsert LM;
                                                                                                                                                                                                                                                                                     (ISIS-) ISIS PHARM INC. (ABBO ) ABBOTT LAB.
                                                                                                                                                                                                                                                                                                                                      WPI; 2002-519883/55.
                                                                                                                                                                                                       WO200250248-A2
                                                                                                                       modified base
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 modified_base
                              modified_base
                                                           modified_base
                                                                                        modified_base
                                                                                                                                                               modified_base
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/*tag= g /mod_base= m5c /*tag= c /mod_base= OTHER /mod_base= OTHER 19. .20

/*tag= h /mod_base= m5c

ч

'note= "2-methoxyethyl (2'-MOE) nucleotides"

mod_base= OTHER

*tag= b

/*tag= d /mod_base= m5c

/*tag= e /mod_base= m5c

/*tag= f /mod_base= m5c

/*tag= a /mod_base= OTHER /not_e= "Phosphorothioate backbone"

Location/Qualifiers

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The invention relates to antisense compounds targetted to a nucleic acid encoding human receptor interacting protein (RIP)2 to inhibit its expression. Antisense compounds are used for treating diseases associated with RIP2 expression. They are also useful in antisense gene therapy. The present sequence is an oligonucleotide targetted to human RIP2 DNA
encoding human receptor interacting protein (RIP)2, for treating diseases associated with RIP2 expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 84.2%; Pred. No. 2.46+02;
Matches 16; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 1 A; 8 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                               Claim 3, Col 46, 35pp; English.
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Gaps

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Query Match 3.3%; Score 14.2; DB 1; Length 20; Best Local Similarity 84.2%; Pred. No. 2.4e+02; Matches 16; Conservative 0; Mismatches 3; Indels

308 CCCCGGGGACCGCGTGCTG 326

CTCCGGGGACTGGGTGCTG 2

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RESULT 121
AAD45181/c
1D AAD45181 standard; DNA; 20 BP.
XX
AC AAD45181;
XX
XX
XY
XY
TOEC-2002 (first entry)

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Gaps

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Anti-human type II DNA topoisomerase alpha antibody-related DNA #38
                                                                                              Human; type II DNA topoisomerase alpha antibody epitope; ss.
                      29-NOV-2002 (first entry)
                                                                                                                                                                                        JP2002191364-A.
                                                                                                                                                                                                                              09-JUL-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABI93857;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABI93857
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes a method (M1) for the diagnosis of dentinogenesis imperfecta type II and/or its accompanying deafness comprising determining the dentin sialophosphoprotein (DSPP) gene, its transcript and/or protein of an individual for comparison of their sequences with the normal equences and judging the individual to have higher risk of suffering from the disease then the normal population. Also described are: (1) treating dentinogenesis imperfecta type III and/or its accompanying deafness by administering a safe and effective dose of normal DSPP and/or DSP protein to patients; (2) drug compositions containing safe doses of DSPP and/or DSP protein; and (3) a regent kit for detecting dentinogenesis imperfecta type II and/or its accompanying deafness containing probes for binding to the mutation site. The DSPP gene and protein sequences have auditory activity. The method (M1), dentin sialophosphoprotein (DSPP) gene and DSP protein are useful for diagnosing and treating imperfecta type II and/or its accompanying deafness. The DSPP gene is located to chromosome 4421. The present sequence represent as PCR primer for the human DSPP gene, which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                     Human, dentin sialophosphoprotein precursor; dentin sialophosphoprotein;
DSPP; dentinogenesis imperfecta type II; deafness; auditory;
chromosome 4q21; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Diagnosis of dentinogenesis imperfecta type III and its accompanying deafness using dentin stalophosphoprotein gene and encoded products.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.3%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 2.4e+02; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 4 A; 3 C; 7 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (SHAN-) SHANGHAI RES CENT BIOTECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hu L;
                                                                                                                                                                                                                 Human DSPP PCR primer SEQ ID NO:15.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 3; Page 12; 38pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Xiao S, Zhao G, Yu C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    353 CTACAGCGACTTCCTCACT 371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
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                                                                                       ABQ73550 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-AUG-2001; 2001WO-CN001292.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-SEP-2000; 2000CN-00125042.
                                                                                                                                                                        03-OCT-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-557897/59.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                     WO200258722-A1.
                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                01-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                           ABQ73550;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kong X,
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                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a target protein fused with a polypeptide having an amino acid sequence containing an epitope of anti-human type II DNA topolsomerase alpha antibody and the DNA encoding it. The sequences can be used in a method for the detection or the determination of a target protein in which the target protein is detected or determined by using the reactivity between the target protein and the above fused protein as the index, and also in a method for the purification of a target protein in which the above fused protein is contacted to anti-human type II DNA topolsomerase alpha antibody carried on a solid carrier. This sequence represents DNA encoding an anti-human type II DNA topolsomerase alpha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human, K-ras, PCR primer, probe, capture probe, mutation detection; ligase detection reaction, LDR, p53; BRCAL; BRCA2; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndrome, obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forenal; environmental monitoring; food industry; feed industry; 88.
                                                                                                                                                                                                          Detection or determination of a protein, a fused protein, a DNA, a vector, purification of a target protein, a solid carrier, an epitope peptide, a kit for the detection or determination.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
3.3%; Score 14.2; DB 1; Length 20;
Best Local Similarity 84.2%; Pred. No. 2.4e+02;
Matches 16; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 3 A; 7 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Capture oligonucleptide Zip ID#944 oligo #9.
                                                                                                                                                                                                                                                                                                                             Disclosure; Page 33; 38pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26 CGAGGGCTGGGACGAGAT 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CGAGAGCTGGGACATAGAT 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP.
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26-DEC-2000; 2000JP-00394675.
                                                    26-DEC-2000; 2000JP-00394675.
                                                                                                        (MITU ) MITSUBISHI CHEM CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABI93857 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                            WPI; 2002-594353/64.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antibody epitope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200179548-A2.
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BB

ABS66287 standard; DNA; 20

RESULT 123
ABS66287/c
ID ABS662
XX
AC ABS662

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ABS66287

Gerry NP, Favis R, Kliman R; (CORR ) CORNELL RES FOUND INC. 14-APR-2000; 2000US-0197271P. WPI; 2002-034366/04. Barany F, Zirvi M, 

Designing capture oligonucleotide probes for use on a support to which complementary oligonucleotides hybridize with little mismatch.

Example 5; Fig 29; 300pp; English.

The present invention describes a method (MI) for designing capture oligonucleotide probes (I) for use on a support to which complementary oligonucleotide probes (II) will hybridise with little mismatch, where (I) have melting temperatures within a narrow range. The method is useful for detecting infectious diseases caused by bacterial infectious agents e.g. Cryptococcus neoformans, Candida albicans and harborial and pollowing and parasitius infectious agents of special man pollowing, wituses e.g. T-cell lymphocytotrophis cirus, bepergillus fumigautus, viruses e.g. T-cell lymphocytotrophis cirus, casteria and pollowirus, Entamoeba histolytica and Dracunculus medinesis. The method is also useful for detecting genetic diseases such as I hydroxylase deficiency, Turner Syndrome and obssity defects.

CC detecting cancer involving oncogenes, tumour suppressor genes, or genes involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCA1 gene, cancer is specifically associated with a gene selected from BRCA1 gene, cancer is specifically associated with a gene selected from BRCA1 gene, cancer is stop and infrared microscope the support at the method is also used for environmental monitoring, forensics and the food and feed industry, detecting comprises scanning electron microscope and infrared microscope) the support at the particular sites and identifying if ligation of the oligomucleotide probe present coligonucleotide sequences. AB182074 to a presence or absence of the target nucleotide sequences. Basence of the target nucleotide sequences. AB182074 to of the present invention

Sequence 20 BP; 2 A; 8 C; 4 G; 6 T; 0 U; 0 Other;

ö 0; Gaps Query Match 3.3%; Score 14.2; DB 1; Length 20; Best Local Similarity 84.2%; Pred. No. 2.46+02; Matches 16; Conservative 0; Mismatches 3; Indels

25 CCGAGGCTGGGACGAAGA 43 20 CCGTGGGATAGGACGAAGA 2 ઠ

ABZ98645 standard; DNA; 20 BP ABZ98645; RESULT 125 ABZ98645/ 

Human tryptase a oligonucleotide sequence. 17-OCT-2003 (first entry)

Human, antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

WO200285308-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013135. 24-APR-2001; 2001US-0286137P

(EPIG-) EPIGENESIS PHARM INC.

Katz E, Pabalan J, Aguilar D; Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Nyce JW, I Miller S,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure, SEQ ID NO 13887; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonuclectide antisense to the intiation coodon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 mucleotides of unctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entiinflammatory steroid and ubjournone. A composition of the invention has antiinflammatory antiallargic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antisflammatory steroid in a subject, for reducing levels of adenosine or lung surfactant in a subject, for reducing levels of adenosine conting antisflammation, lung allergies, or a respiratory disease or condition. Ung inflammation, lung allergies, or a respiratory disease or condition. Specification, but was obtained in electronic format directly from WIPO at the wipo.int/published_pot_sequences

Sequence 20 BP; 3 A; 5 C; 7 G; 5 T; 0 U; 0 Other;

Gaps .; 0 3.3%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 2.4e+02; tive 0; Mismatches 3; Indels Matches 16; Conservative Best Local Similarity Query Match

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RESULT 126 ADE27892,

892/c ADE27892 standard; DNA; 20 BP.

ADE27892;

Human B7-1 targeted oligonucleotide SEQ ID 154.

29-JAN-2004 (first entry)

ss; human; B7-1; inflammatory skin disorder; antisense; psoriasis; contact dermatitis; atopic dermatitis; seborrheic dermatitis; nummular dermatitis; generalised exfoliative dermatitis; eczema; critical costimulatory molecule.

Synthetic

Homo sapiens.

US2003176374-A1.

18-SEP-2003.

09-MAY-2001; 2001US-00851871.

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The invention relates to a method of treating an inflammatory skin disorder in an individual by topically applying an antisense compound targeted to a nucleic acid molecule encoding a human B7 protein. The invention is for treating an inflammatory skin disorder in individual. The skin disorder is psoriasis, contact dermatitis, atopic dermatitis, seborrheic dermatitis, nummular dermatitis, generalised exfoliative dermatitis or eczema. The invention effectively modulates critical costinulatory molecules such as the B7 protein. The present sequence represents a human B7-1 targeted oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Isolated DNA molecule encoding cholecystokinin receptor protein - are
                                                                                                                                                                              Treating an inflammatory skin disorder such as psoriasis comprises topically applying an antisense compound targeted to the nucleic acid encoding human B7 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cholecystokinin receptor protein; CCK; gastrointestinal receptor; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence of nested PCR primer for cholecystokinin (CCK) cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                               3.3%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 2.4e+02; tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 6 A; 8 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                           Example 12; SEQ ID NO 154; 88pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DEPT HEALTH & HUMAN SERVICE
                                                                                                                            Karras JG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             398 GAAGGICIICIACGIGAIC 416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19 GAAGGTGTTCTTCGTGAGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         92US-00831248.
92US-00861769.
92US-00928033.
92US-00937609.
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            31-DEC-1996; 96US-00777266.
04-JUN-1999; 99US-00326186.
25-MAY-2000; 2000WO-US014471.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1676/c
AAQ47676 standard; cDNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (revised)
(first entry)
                                                                                                                            Bennett CF, Vickers TA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity 84.2
nes 16; Conservative
                                                                   (BENN/) BENNETT C F.
(VICK/) VICKERS T A.
(KARR/) KARRAS J G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (PI; 1993-272886/34.
                                                                                                                                                        WPI; 2003-863863/80.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-APR-1992;
11-AUG-1992;
02-SEP-1992;
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07-FEB-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28-JAN-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SU (HSSU)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAQ47676,
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                                                                                       Mixed oligos primed amplification of CCK cDNA was performed using 2 groups of degenerate primers based on the AA sequence from AAR38890. The sense gp. of primers was 72 fold degenerate (AAQ47672). The anti- gp. of primers was 80 fold degenerate and consisted of AAQ47673. The atti- gp. of primers was 80 fold degenerate and consisted of AAQ47673 & AAQ47674. The product of the PCR was used to generate nondegenerate primers for subsequent PCR. The remaining 3' coding and UTRS was obefor using amplificn. (RACE) of CDNA and anchored PCR. RACE was performed using AAQ47675 for the first round and a nested primer, AAQ47676, for the second round. Anchored PCR used the gene specific primer AAQ47677 and the sequences was cloned using PCR. The sense primer was AAQ47677 and the sequences was cloned using PCR. The sense primer was AAQ47679 and the antisense primer was AAQ47680. (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New nucleic acid segment comprising one of the 10 - 100 bp sequences given in the specification (sequences of a polymorphic site), or the complement of the segment and a method of analysing a nucleic acid comprising determining the base occupying the polymorphic fixe of the polymorphic fixagment sequences are disclosed in the specification. The information obtained from nucleic acid analysis by the method described is useful in diagnosis or monitoring of diseases like cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New nucleic acid segments containing polymorphic sites, or complements and methods of detecting a nucleic acid - for general use including diagnosis and monitoring of diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
purified to isolate cholecystokinin receptor clones and produce anti-
cholecystokinin receptor antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ss; polymorphic site; nucleic acid analysis; diagnosis; monitoring; cancer; inflammation; heart disease; CNS disease.
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                                                                                                                                                                                                                                                                                                                                                                                                       Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleotide fragment containing polymorphic site, WI-7038.
                                                                                                                                                                                                                                                                                                                                                                                                                                               3; Indels
                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 6 A; 7 C; 7 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                  7. Match 3.3%; Score 14.2; DB 1; Local Similarity 84.2%; Pred. No. 2.7e+02; LB 16; Conservative 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Berno A;
                                                             Example; Page 38; 110pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      241 GCTGCTTCCCGGGCTCGGC 259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 10; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20 GCTGCTGCCAGTGCTCGGC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fan J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAV67403 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      98WO-US004571
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97US-0042125P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (AFFY-) AFFYMETRIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chee M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1998-495419/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-DEC-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-MAR-1997;
28-MAR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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Gaps ö |||||||| TGGAGGCAAGGTTCGACTG

21

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AAZ25089 standard; DNA; 21

AAZ25089;

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This sequence is an inhibitor of the human AUR2 protein of the invention. The AUR1 and AUR2 proteins can be used to identify specific modulators of, and to generate specific antibodies recognising AUR1 and AUR2. The modulators can be used for treating conditions involving abnormal AUR signal transduction, specifically cancer (of colon, breast, Kidhey, chandrosarcoma and panneratic tumours, particularly of colon (specifically), breast or kidhey). The modulators can also be used for studying their effects in animal models of proliferative disease. Probes, based on the coding sequences are used, diagnosfically, to detect or quantify AUR mRNA, by hybridisation or polymerase chain reaction (PCR). The DNA, optionally mutated, are useful in gene therapy. Ab are used as diagnostic immunoassay reagents for detecting the proteins
                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New nucleic acid encoding human AUR1 and 2 polypeptides, used to identify specific modulators for treating cancer or for diagnosis.
inflammation, heart disease, CNS diseases, and susceptibility to infection by microorganisms. In addition, the nucleic acid segments are useful in manufacturing medication in the treatment of prophylaxis of diseases, and also the use of the DNA segments as pharmaceutical
                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AUR1; AUR2; human; AUR modulator; cancer; glioma; medullablastoma; chondrosarcoma; pancreatic tumour; proliferative disease; diagnosis; therapy; inhibitor; ss.
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o
                                                                                                                        Query Match 3.3%; Score 14.2; DB 1; Length 21; Best Local Similarity 76.2%; Pred. No. 2.7e+02; Matches 16; Conservative 1; Mismatches 4; Indels
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                                                                                         Sequence 21 BP; 3 A; 4 C; 7 G; 6 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 24; Page 120; 153pp; English.
                                                                                                                                                                                                      GCCACCACTCAGAGGAGTCTC 66
                                                                                                                                                                                                                         21 GCCATCACGCRGAAAGTCTC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99WO-US001283
                                                                                                                                                                                                                                                                                                                              AAX99728 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                      29-SEP-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Plowman GD, Mossie K;
                                                                                                                                                                                                                                                                                                                                                                                                                                       Human AUR2 inhibitor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1999-458699/38.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (SUGE-) SUGEN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9937788-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                   AAX99728;
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                                                                                                                                                                                                                                                                                         RESULT 12:
AAX99728/c
ID AAX99
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3.3%; Score 14.2; DB 1; Length 21; llarity 84.2%; Pred. No. 2.7e+02; Conservative 0; Mismatches 3; Indels

Query Match Best Local Similarity Matches 16; Conservat

148 TGGAGGCCGGCTTCGACTG 166

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The present invention describes human mitogen-activated protein kinase/extracellular response kinase (MAPK/ERK) kinase kinase (MEKK).

specifically designated MEKKI, MEKKZ and MEKKI. The MEKK proteins are used to modulate and regulate signal transduction in cells, as well as for regulation of gene transcription in a cell encoding MEKK, where the call is involved in inflammation, regulation of cellular proliferation and differentiation, regulation of development, regulation of cell death or regulation of the present also used to prepare antibodies. MEKK polymucleorides can be used to produce the protein recombinantly and as a source of probe and primers. The present sequence represents a PCR primer for human MEKK2, which is used in an example from the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New human MEKK polynucleotides and polypeptides, used for regulating signal transduction in cells.
                                                                                                          MEKKI; MEKK2; MEKK3; mitogen-activated protein kinase; MAPK; ERK; extracellular regulated kinase; signal transduction; regulation; MAPK/ERK; MEK; MKK; inflammation; cellular proliferation; differentiation; development; cell death; PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
3.3%; Score 14.2; DB 1; Length 21;
Best Local Similarity 84.2%; Pred. No. 2.7e+02;
Matches 16; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21 BP; 5 A; 3 C; 11 G; 2 T; 0 U; 0 Other;
                                                                              Human MEKK2 PCR primer SEQ ID NO:28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 2; Page 64; 159pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         289 AGCTGGTGAAGGACCTGAG 307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21
                                                                                                                                                                                                                                                                                                                                       98US-0078153P.
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                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                       (CADU-) CADUS PHARM CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1999-571843/48.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18-SEP-2000
                                               09-DEC-1999
                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                          WO9947686-A2
                                                                                                                                                                                                                                                                                                        15-MAR-1999;
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04-SEP-1998;
                                                                                                                                                                                                                                                                         23-SEP-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                       Johnson GL;
                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAA52302;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAA52302
ID AAA5
XX
AC AAA5
XX
DT 18-S
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69

Page

Oligonucleotide used to construct UpEt-Ubi vector, SEQ ID NO:31.

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Plasminogen, human, kringle 5 domain, endothelial cell proliferation, and,ogenesis, antiproliferative, antiarteriosclerotic; cytostatic; antiarfilammatory; antiulcer; antirheumatic; antiarthritic; antiangiogenic; cancer; tumour; autodammune disease; Escherichia coll; recombinant expression; vector construction; PCR primer; se.
                                                                                                                                                                                                                            Preparation of Kringle five peptide fragment for treating various disorders such as angiogenic, ocular, skin diseases and cancer, involves mixing mammalian plasminogen and elastase followed by incubation and isolation.
                                                                                                                                                    96US-00643219.
                                                                                                                                    97US-00851350
                                                                                                                                                                                                                  WPI; 2000-349573/30.
                                                                                                                                                                               (ABBO ) ABBOTT
                                                                                                                                    05-MAY-1997;
                                                                                                                                                    03-MAY-1996;
03-APR-1997;
                                                                                                                                                                                                  Davidson DJ;
                                                                                                JS6057122-A
                                                                                                                   02-MAY-2000
                                                                               Synthetic.
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The invention relates to a method of preparing plasminogen kringle 5

peptide fragments. The method comprises mixing mammalian plasminogen and
elastase in the ratio 1:100-1:300, followed by incubating and isolating
the fragment. The kringle 5 peptides are inhibitors of angiogenesis and
endothelial cell proliferation and migration. The peptides are useful for
treating angiogenic diseases, primary and metastatic solid tumours and
carcinomas of various organs such as breast, genital tract, endocrine
glands, skin, tumours of the brain and eyes and solid tumours arising
creation used for the prophylaxis of various autoimmune diseases (e.g.,
from haematopoietic malignancies such as leukaemias and lymphomas. They
are also used for the prophylaxis of various autoimmune diseases (e.g.,
blood vessel diseases (e.g., haemangiomas, Osler-Webber Syndrome),
creamiated by excessive or abnormal stimulation of Endothelial cells
(e.g., Crohn's disease, atherosclerosis), diseases are also useful as a birth control agent which
inhibits ovulation and establishment of the placenta. Sequences AAA52294AS2304 represent PCR primers used in the construction of Escherichia coll
expression vectors for recombinant expression of various human Example 20; Col 49; 48pp; English plasminogen kringle 5 fragments

Sequence 21 BP; 7 A; 6 C; 8 G; 0 T; 0 U; 0 Other;

Gaps ö 3.3%; Score 14.2; DB 1; Length 21; 84.2%; Pred. No. 2.7e+02; ive 0; Mismatches 3; Indels Query Match Best Local Similarity 84.2 Matches 16; Conservative

380 CCGCGACGACGCCCAAG 398

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CGGCGACGACGACAAG 21 ო 셤

AAA52303 standard; DNA; 21 (first entry) 18-SEP-2000 AAA52303; RESULT 132 AAA52303 KAKKAK

Oligonucleotide used to construct UpEt-Ubi vector, SEQ ID NO:32

Plasminogen, human, kringle 5 domain; endothelial cell proliferation; angiogenssis; antiproliferative; antiarteriosolerotic; cytostatic; antiafficantic; antiafficmatory; antiulcer; antirheumatic; antiarthritic; antiangiogenic; cancer; tumour; autoimmune disease; Escherichia coli; recombinant expression; vector construction; PCR primer; ss.

Synthetic.

US6057122-A.

02-MAY-2000

97US-00851350 05-MAY-1997;

96US-00643219. 97US-00832087. 03-MAY-1996; 03-APR-1997; 

LAB. (ABBO ) ABBOTT

Davidson DJ;

WPI; 2000-349573/30.

Preparation of Kringle five peptide fragment for treating various disorders such as anglogenic, ocular, skin diseases and cancer, involves mixing mammalian plasminogen and elastase followed by incubation and isolation

Example 20; Col 49; 48pp; English.

The invention relates to a method of preparing plasminogen kringle 5 peptide fragments. The method comprises mixing mammalian plasminogen and elastic iiio-iiio0, iollowed by incubating and isolating the fragment. The kringle 5 peptides are inhibitors of angiogenesis and endothelial cell proliferation and migration. The peptides are useful for treating angiogenic diseases, primary and metastatic solid tumours and carcinomas of various such as breast, gental tract, endocrine glands, skin, tumours of the brain and eyes and solid tumours arising from haematopoietic malignancies such as leukaemias and lymphomae. They are also used for the prophylaxis of various autoimmune diseases (e.g., psortasis), blood vessel diseases (e.g., haemangiomas, Osler-Webber Syndrome), cliseases caused by excessive or abnormal stimulation of endothelial cells (e.g., Crohn's disease, atherosclerosis), diseases which have angiogenesis as a pathologic consequence (e.g., cat scratch disease and ulcren). The peptides are also useful as a birth control agent which and establishment of the placents. Sequences AAA52294. A52304 represent PCR primers used in the construction of Escherichia coli expression vectors for recombinant expression of various human fragments plasminogen kringle 5

Sequence 21 BP; 0 A; 8 C; 6 G; 7 T; 0 U; 0 Other;

Gaps ö 3.3%; Score 14.2; DB 1; Length 21; 84.2%; Pred. No. 2.7e+02; tive 0; Mismatches 3; Indel8 Local Similarity 84.2 les 16; Conservative Query Match Best Loca Matches

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AAF29947 standard; DNA; 21 AAF29947; RESULT 133
AAF29947/c
XX
AC AAF299
XX
AC AAF299
XX
XX
DE Primer

Вb

(first entry) 05-APR-2001

Primer #5

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BP.
                                                                                10-SEP-1999; 99US-0153357P.
26-JUL-2000; 2000US-0220947P.
16-AUG-2000; 2000US-0225724P.
                                                      07-SEP-2000; 2000WO-US024503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAF97092 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16; Conservative
                                                                                                                                                                                     Lander ES, Gargill M,
                                                                                                                                                                                                                   WPI; 2001-226749/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200118250-A2
WO200118250-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-JJN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-MAR-2001
                           15-MAR-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Key
Variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAF97092;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ò
                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to a cholecystokinin (CCK) receptor protein. The CCK receptor-encoding DNA molecule is useful for expressing and purifying CCK receptor protein to sequenceable-grade homogeneity. The CCK receptor proteins or fragments are useful for obtaining antibodies that can recognize CCK-expressing cells. The transformed eukaryotic cell innes are useful for studying the receptor in an environment similar to its native environment, e.g. in the context of studying the electrophysiology or binding properties of the receptor. The transformed producel large amounts of the receptor for immunological purposes or for studying protein structure, e.g. crystallography
                                                                                                                                                                                                                                                                                                                       New cholecystokinin (CCK) receptor-encoding DNA molecule, useful for
producing and purifying human CCK receptor protein to sequenceable-grade
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; variant thrombospondin 1; variant thrombospondin 4; SNP; polymorphism; vascular disease; coronary artery disease; forensics; mycoardial infarction; atherosclerosis; stroke; venous thromboembolism; pulmonary embolism; paternity test; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /*tag= a
/standard_name= "single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ..
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ouery Match 3.3%; Score 14.2; DB 1; Length 21; Best Local Similarity 84.2%; Pred. No. 2.7e+02; Matches 16; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 6 A; 7 C; 7 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human gene single nucleotide polymorphism #895.
                 Cholecystokinin; CCK receptor; purify; ss.
                                                                                                                                                                                                                                    (USSH ) US DEPT HEALTH & HUMAN SERVICES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Col 11; 82pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           241 GCTGCTTCCCGGGCTCGGC 259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20 GCTGCTGCCAGTGCTCGGC 2
                                                                                                                                                           92US-00831248.
92US-00861769.
92US-00928033.
92US-00937609.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAF96134 standard; DNA; 21 BP.
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                                                                                                                               93US-00029170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                            WPI; 2001-136725/14.
                                                                                                                                                                         01-APR-1992;
11-AUG-1992;
02-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-JUN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                             Unidentified
                                                                        US6169173-B1
                                                                                                                                  10-MAR-1993;
                                                                                                                                                                                                                                                                                                                                       producing an homogeneity.
                                                                                                    02-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Key
Variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAF96134;
                                                                                                                                                                                                                                                                Wank SA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAF96134

YA AAF9

AAC AAF9

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YX YA

DT XX

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XX HOME

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XX YA

YA YET

YA YA YA

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The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various opportments of polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, concary heart disease, stroke, peripheral vascular diseases, venous thromboembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human, variant thrombospondin 1, variant thrombospondin 4, SNP,
polymorphism, vascular disease, coronary artery disease, forensics,
myocardial infarction, atherosclerosis, stroke, venous thromboembolism,
pulmonary embolism, paternity test, ds.
                                                                                                            Mccarthy JJ;
                                                                                                                                                                                                                                            Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and atherosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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84.2%; Pred. No. 2.7e+02;
iive 0; Mismatches 3; Indels
                                                                                                            Daley GQ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 1 A; 8 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human gene single nucleotide polymorphism #1853.
                                                                                                        Ireland JS, Bolk S,
RES.
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replace(11,C)
/*tag= a
(WHED ) WHITEHEAD INST BIOMEDICAL (MILL-) MILLENNIUM PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example; Page 111; 242pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CTGGCCCGCCTGGCGGTGG 150
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16-AUG-2000; 2000US-0225724P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO2003013534-A2.
                                                                                                                                 atherosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Heinrich G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-FEB-2003
                                                      Lander ES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ACF62200;
                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 137
                                                                                                                                                                                                                                                                                                                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                8
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                                                                                                                                                                                                                  The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various bolymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thromboembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; variant thrombospondin 1; variant thrombospondin 4; SNP;
polymorphism; vascular disease; coronary artery disease; forenaics;
myocardial infarction; atherosclerosis; stroke; venous thromboembolism;
pulmonary embolism; paternity test; ds.
                                                                                                  Mccarthy JJ;
                                                                                                                                          Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and
                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /*tag= a
/standard_name= "single nucleotide polymorphism"
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                                                                                                                                                                                                                                                                                                                                                                           3.3%; Score 14.2; DB 1; Length 21; 84.2%; Pred. No. 2.7e+02; ive 0; Mismatches 3; Indels
                                                                                                  Daley GQ,
                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 4 A; 7 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human gene single nucleotide polymorphism #2100.
                                                                                                  o,
                                                                                                Bolk
                                                                (WHED ) WHITEHEAD INST BIOMEDICAL RES (MILL-) MILLENNIUM PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
replace(11,T)
                                                                                                  Ireland JS,
                                                                                                                                                                                                 Example; Page 174; 242pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                       GGCGCCACCAAGCTGGTGA 297
                                                                                                                                                                                                                                                                                                                                                                                                                                     GGTGGCACAAAGCTGATGA 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAF97339 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-SEP-1999; 99US-0153357P.
26-JUL-2000; 2000US-0220947P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-SEP-2000; 2000WO-US024503
                      10-SEP-1999; 99US-0153357P.
26-JUL-2000; 2000US-0220947P.
16-AUG-2000; 2000US-0225724P.
  07-SEP-2000; 2000WO-US024503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-JUN-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                Gargill M,
                                                                                                                     WPI; 2001-226749/23
                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
ses 16; Conserv
                                                                                                                                                                            atherosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200118250-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-MAR-2001
                                                                                                  ES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Key
Variation
                                                                                                                                                                                                                                                                                                                                                                                                                       279
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAF97339;
                                                                                                                                                                                                                                                                                                                                                                                                                                           21
                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                  Lander
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 136
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p450,
                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various opportantly objective within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thromboembolism pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
                                                                                           Mccarthy JJ
                                                                                                                                                                                                                  Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cancer; CYP3A5; irinotecan; pharmaceutical; malignant glioma;
cytochrome p450; subfamily IIIA; nifedipine oxidase; polypeptide
cytostatic; PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 14.2; DB 1; Length 21;
Pred. No. 2.7e+02;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cancer based on CYP3A5 related oligonucleotide SEQ ID NO:1.
                                                                                              Daley GQ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21 BP; 4 A; 3 C; 11 G; 3 T; 0 U; 0 Other;
                                                                                              Bolk S,
(WHED ) WHITEHEAD INST BIOMEDICAL RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
                                                                                              Ireland JS,
                                                                                                                                                                                                                                                                                                                                                                                    Example, Page 192; 242pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17 GCGGGTGACCGAGGGCTGG 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3 GTGGGTGACCCAGGGTGG 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.3%;
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24-MAY-2002; 2002EP-00011710.
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                                 (MILL-) MILLENNIUM PHARM INC
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hes 16; Conservative
                                                                                           Gargill M,
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Disclosure, Page 32, 86pp; English.
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The present invention describes the use of irinotecan (1) or its derivative for the preparation of a pharmaceutical composition for treating colorectal, gastric, lung, ovarian or pancreatic cancer, or malignant gliona in a subject having a genome with a variant allele which comprises a cytochrome p450, subfamily IIIA (nifediplne oxidase), polypeptide 5 (CYPAAS) polymuclectide (II). (I) and (II) have cytostatic activity. The therapeutic applications of (I) is improved, since it is possible to individually treat a subject with an appropriate consequency and effects are efficiently avoided. Unnecessary and potentially harmful treatment of those subjects who do not respond to the treatment with mubstrances (nonresponders), as well as the development of treatment with understances (nonresponders), as well as the development of charge resistances due to suboptimal drug dosing can be avoided, ACF62200 to ACF62751 and ABM34912 to ABM36013 represent sequences used in the exemplification of the present invention

Sequence 21 BP; 0 A; 6 C; 9 G; 6 T; 0 U; 0 Other;

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Gapa
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3.3%; Score 14.2; DB 1; Length 21; 84.2%; Pred. No. 2.7e+02; tive 0; Mismatches 3; Indels
                                                              336 GACCAGGGCCGGCTGCTCT 354
                                                                                 16, Conservative
                 Best Local Similarity
Matches 16; Conserv
      Query Match
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Cancer based on CYP3A5 related oligonucleotide SEQ ID NO:2. ACF62201 standard; DNA; 21 (first entry) 08-OCT-2003 ACF62201; RESULT 138 

BP.

Cancer; CYP3A5; irinotecan; pharmaceutical; malignant glioma; cytochrome p450; subfamily IIIA; nifedipine oxidase; polypeptide cytostatic; PCR primer; ss.

5

Synthetic.

WO2003013534-A2.

20-FEB-2003.

23-JUL-2002; 2002WO-EP008219.

23-JUL-2001; 2001EP-00117608. 24-MAY-2002; 2002EP-00011710.

(EPID-) EPIDAUROS BIOTECHNOLOGIE AG.

Heinrich G, Kerb R;

WPI; 2003-268144/26.

New use of irinotecan for preparation of compositions for treating in subject having genome with variant allele comprising cytochrome subfamily IIIA, polypeptide 5 polynucleotide, termed CYP3A5.

Disclosure; Page 32; 86pp; English.

The present invention describes the use of irinotecan (I) or its derivative for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or panoreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a cytochrome p450, subfamily IIIA (infedipine oxidase), polypeptide 5 (CYP3A5) polynucleotide (II). (I) and (II) have

Sequence 21 BP; 6 A; 9 C; 6 G; 0 T; 0 U; 0 Other;

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The present invention describes a method for the use of irinotecan (I) or its derivative for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancratic cancer, or malignant gliona in a subject having a genome with a variant allele which comprises a multidrug resistance protein 1 (MRP1) allele which comprises a multidrug resistance protein 1 (MRP1) can be used for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant gliona in a subject, where the subject is a human (preferably African or Asian) or a mouse. The present sequence represents a sequence which is used in the exemplification of the present invention.
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cytostatic activity. The therapeutic applications of (I) is improved, since it is possible to individually treat a subject with an appropriate decivative of (I). Therefore, undesirable, harmful or toxic effects are efficiently avoided. Unnecessary and potentially harmful treatment of those subjects who do not respond to the treatment with substances (nonresponders), as well as the development of drug resistances due to suboptimal drug dosing can be avoided. ACF62200 to ACF62751 and ABM34912 to ABM35013 represent sequences used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Use of irinotecan or its derivative for preparation of a pharmaceutical composition for treating cancer in a subject having a genome with a variant allele comprising a multidrug resistance protein 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               irinotecan; colorectal cancer; cervical cancer; gastric cancer;
lung cancer; ovarian cancer; pancreatic cancer; malignant glioma;
variant allele; multidrug resistance protein 1; MRP1; cytostatic; gene;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MRP1 based cancer related nucleic acid SEQ ID NO:2.
                                                                                                                                                                                                Sequence 21 BP; 6 A; 9 C; 6 G; 0 T; 0 U; 0 Other;
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24-MAY-2002; 2002EP-00011710.
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New isolated DNA molecule encoding a cholecystokinin (CCK) receptor protein, useful for neuroendocrine modulation of the immune system, and for obtaining antibodies that can recognize CCK-expressing cells.
                                                                                                 Rat; ss; PCR; primer; CCK; cholecystokinin receptor; immunomodulator; RACE: rapid amplification of cDNA ends.
                                                                        RACE oligonucleotide #2 used to amplify rat CCK cDNA 3' sequences.
 ACD26205 standard; DNA; 21 BP
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                                                                                                                                                                    US2003055238-A1.
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02-SEP-1992;
10-MAR-1993;
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                                                   13-SEP-2003
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                                                                                                                                            Rattus sp.
                          ACD26205;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Use of irinotecan or its derivative for preparation of a pharmaceutical composition for treating cancer in a subject having a genome with a variant allele comprising a multidrug resistance protein 1 polynucleotide.
                                                                                                                                                                                                                                                  irinotecan; colorectal cancer; cervical cancer; gastric cancer;
lung cancer; ovarian cancer; pancreatic cancer; malignant glioma;
variant allele; multidrug resistance protein 1; MRP1; cytostatic; gene;
                             Gaps
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34.2%; Pred. No. 2.7e+02;
Ive 0; Mismatches 3; Indels
Score 14.2; DB 1; Length 21;
Pred. No. 2.7e+02;
0; Mismatches 3; Indels
                                                                                                                                                                                                                          WRP1 based cancer related nucleic acid SEQ ID NO:1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
                                                      336 GACCAGGGCCGGCTCT 354
                                                                                                                                               BP
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24-MAY-2002; 2002EP-00011710.
3.3%;
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Best Local Similarity 84.2%;
Matches 16; Conservative
                                                                     ADB20871 standard; DNA; 21
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Query Match
Best Local Similarity 84.2'
Matches 16; Conservative
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92US-00831248. 92US-00861769. 92US-00928033. 92US-00937609.

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99US-00443745

(first entry)

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                                             This invention relates to a novel isolated DNA molecule encoding a cholecystokinin (CCK) receptor protein. The invention also discloses a method for purifying a CCK receptor in 1% digitionin, applying the elucation containing CCK receptor in 1% digitionin, applying the solubilised receptor preparation to a cationic exchange resin and purifying the elucate of the resin. The purified elucate is then added to an agarose-bound lectin and applied the elucate to a cabacron blue sepharose column and a CCK receptor protein of sequenceable-grade purity. The CCK receptor protein of the invention may have immunomidulatory activity. The DNA molecule of the invention may have immunomidulatory activity. The DNA molecule of the invention is useful for purifying CCK receptor protein modulation of the immune system, and for a setuincendocrine modulation of the immune system, and for obtaining antibodies that can recognise CCK-expressing cells. The present sequence represents a RACE PCR PRIMET used to amplify the 3, end of the invention of the immune system.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cholecystokinin (CCK) receptor cDNA sequence of the invention
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Example 1; Page 6; 83pp; English
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336 GACCAGGGCCGCTGCTCT 354 Greengecegerer 19

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RESULT

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Query Match
Best Local Similarity 84.2
Matches 16; Conservative
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20-FEB-2003
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ADB96944/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Use of irinotecan to treat cancer patient by determining if patient has variant alleles of UGT1A1 gene, administering increased/decreased amounts of irinotecan based on increased/decreased levels of UGT1A1 gene product.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ss; irinotecan; cancer; UGT1A1; cytostatic; topoisomerase I inhibitor; colorectal cancer; cervical cancer; gastric cancer; lung cancer; ovarian cancer; pancreatic cancer; malignant glioma; uridine diphosphate glycosyltransferasel member A1.
                                  ss; irinotecan; cancer; UGTIA1; cytostatic; topoisomerase I inhibitor; colocectal cancer; cervical cancer; astric cancer; lung cancer; ovarian cancer; pancreatic cancer; malignant glioma; uridine diphosphate glycosyltransferasel member Al.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human UGT1A1 variant allele sequence fragment SEQ ID NO:1.
Human UGT1A1 variant allele sequence fragment SEQ ID NO:2.
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                                                                                                                                                                                                                                                                                                                                     (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
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24-MAY-2002; 2002EP-00011710.
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                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                        20-FEB-2003.
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8X8686666666668X8X171728X1X3X3X8X8X8X8X8X8

ADB87960;

RESULT 143 ADB87960

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The invention relates to the novel use of irinotecan to treat a patient suffering from cancer. This involves determining if the patient has one or more variant allales of the UGTAL gene, and if the patient has one or more of such variant allales, irinotecan is administered in an increased or decreased amount in comparison to the amount that is administered without regard to the patient's allales in the UGTALA gene. The invention has cytostatic activity. A composition of the invention acts as a topolsomerase I inhibitor. The method is useful for treating a patient, an animal e.g. mouse or a human, preferably African or Asian, suffering from cancer such as colorectal, cervical, gastric cancer, lung, ovarian, pancreatic cancer or malignant glions. The present sequence is udes in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Use of irinotecan to treat cancer patient by determining if patient has variant alleles of UGTIA1 gene, administering increased/decreased amounts of irinotecan based on increased/decreased levels of UGTIA1 gene product.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.3%; Score 14.2; DB 1; Length 21; 84.2%; Pred. No. 2.7e+02; tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human UGT1A1 variant allele sequence fragment SEQ ID NO:2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21 BP; 0 A; 6 C; 9 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
                                                                                                                                                                                                                    (EPID-) EPIDAUROS BIOTECHNOLOGIE AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 8; Page 44; 107pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   336 GACCAGGCCGGCTGCTCT 354
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADB96944 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-JUL-2002; 2002WO-EP008218.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-JUL-2001; 2001EP-00117608.
24-MAY-2002; 2002EP-00011710.
23-JUL-2002; 2002WO-EP008217
                                                                                23-JUL-2001; 2001EP-00117608.
24-MAY-2002; 2002EP-00011710.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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wed Apr 41 14:38:41 4004

WPI; 2003-268145/26.

New use of irinotecan for preparation of pharmaceutical compositions for treating cancer in subject having genome with variant allele comprising multidrug resistance 1 polynucleotide.

Claim 4; Page 69; 130pp; English.

The invention relates to the novel use of irinotecan or its derivative for the preparation of pharmaceutical compositions for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a multidrug resistance 1 (WDR1) polymolectide. A composition of the invention has cytostatic activity. The invention is useful for the preparation of pharmaceutical compositions for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject (preferably human, more preferably African or Asian) or a mouse. The present sequence is used in the exemplification of the

Sequence 21 BP; 6 A; 9 C; 6 G; 0 T; 0 U; 0 Other;

ô 3.3%; Score 14.2; DB 1; Length 21; 84.2%; Pred. No. 2.7e+02; ive 0; Mismatches 3; Indels Query Match
Best Local Similarity 84.2
Matches 16; Conservative

336 GACCAGGGCCGGCTGCTCT 354 ٣ 21 Greenedecedererer ò 셤

ADB96943 standard; DNA; 21 BP RESULT 145 ADB96943

ADB96943;

04-DEC-2003 (first entry)

Human UGT1A1 variant allele sequence fragment SEQ ID NO:1.

irinotecan; colorectal cancer; cervical cancer; gastric cancer; lung cancer; ovarian cancer; panoreatic cancer; malignant glioma; multidrug resistance 1; MDR1; cytostatic; human; ds; Cyp3A5; MRP1; MDR1; TOP1.

Homo sapiens.

WO2003013537-A2.

20-FEB-2003

23-JJL-2002; 2002WO-EP008218.

23-JUL-2001; 2001EP-00117608. 24-MAY-2002; 2002EP-00011710.

(EPID-) EPIDAUROS BIOTECHNOLOGIE AG.

Kerb R; Heinrich G,

WPI; 2003-268145/26.

New use of irinotecan for preparation of pharmaceutical compositions for treating cancer in subject having genome with variant allele comprising multidrug resistance 1 polymucleotide.

Claim 4; Page 69; 130pp; English.

The invention relates to the novel use of irinotecan or its derivative for the preparation of pharmaceutical compositions for treating colorectal cervical, gastric, lung, ovarian or pancreatic cancer, amalignant glioma in a subject having a genome with a variant allele which 

ô comprises a multidrug resistance 1 (MDR1) polynucleotide. A composition of the invention has cytostatic activity. The invention is useful for the preparation of pharmaceutical compositions for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject (preferably human, more preferably African or Asian) or a mouse. The present sequence is used in the exemplification of the Gaps ö Length 21; 3; Indels Sequence 21 BP; 0 A; 6 C; 9 G; 6 T; 0 U; 0 Other; Score 14.2; DB 1; Pred. No. 2.7e+02; 0; Mismatches 3; 336 GACCAGGCCGGCTGCTCT 354 1.3%; Local Similarity 84.2%; 16; Conservative invention. Query Match Matches 888888888 ઠે

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ADB92134

ADB92134 standard; DNA; 21

ADB92134;

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(first entry) 04-DEC-2003 Human UGT1A1 variant allele sequence fragment SEQ ID NO:1.

irinotecan; colorectal cancer; cervical cancer; gastric cancer; lung cancer; ovarian cancer; pancreatic cancer; malignant glioma; multidrug resistance 1; MDR1; cytostatic; ds; human; UGT1A1; MRP1; TOP1.

Homo sapiens,

WO2003013535-A2.

20-FEB-2003.

23-JUL-2002; 2002WO-EP008220

23-JUL-2001; 2001EP-00117608. 24-MAY-2002; 2002EP-00011710.

(EPID-) EPIDAUROS BIOTECHNOLOGIE AG.

Heinrich G, Kerb R;

WPI; 2003-342400/32.

New use of irinotecan for preparation of pharmaceutical compositions for treating cancer in subject having genome with variant allele comprising multidrug resistance 1 polynucleotide.

Disclosure; Page 41; 104pp; English.

The invention relates ro a novel use of irinotecan or its derivative for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, covarian or pancreatic cancer, or malignant gliome in a subject having a genome with a variant allele which comprises a multidrug resistance | (MDR1) polymucleotide. A composition of the invention has cytostatic activity. The present sequence is used in the exemplification of the invention. 

Sequence 21 BP; 0 A; 6 C; 9 G; 6 T; 0 U; 0 Other;

Gaps ö Query Match 3.3%; Score 14.2; DB 1; Length 21; Best Local Similarity 84.2%; Pred. No. 2.7e+02; Matches 16; Conservative 0; Mismatches 3; Indels

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336 GACCAGGCCGGCTGCTCT 354

à g

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The invention relates to an isolated or purified polymucleotide encoding a polypeptide (the wild-type form of which is involved in synaptogenesis) that includes at least one mutation associated with development of eneurological disease and/or a predisposition to development of mental content that modulate their activity. Also nucleic acid, polypeptide, polypeptide are used to screen for agents that modulate their activity. Also nucleic acid, polypeptide, or polypeptide are used to screen for containing the vector, are useful as pharmaceuticals for treating mental and neurological disorders, specifically autism, Asperger containing the nucleic acid and containing the nucleic acid and polypeptide can be used to similarly. Also detecting mutations in the nucleic acid and polypeptide, can be used to detect that affect formation of synapses and to disquose the nucleic formation of synapses and to disquose mental disease. This sequence corresponds to a PCR primer used to amplify the human wild type HNL4X (ADC24764) and HNL4Y (ADC24764) genes.
                                                                                                                                                                                                                                                                                                                                New nucleic acid encoding mutant protein involved in synaptogenesis, useful for treatment and diagnosis of e.g. autism, Asperger syndrome, and schizophrenia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 3 A; 5 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                          (INSP ) INSERM INST NAT SANTE & RECH MEDICALE.
(INSP ) INST PASTEUR.
(ASSI-) ASSISTANCE PUBLIQUE HOPITAUX PARIS.
                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; SEQ ID NO 21; 416pp; French.
                                                                                                                                         30-NOV-2001; 2001CA-02364106
                                                                                                        02-DEC-2002; 2002WO-FR004134
                                                                                                                                                                                                                                                                                                       WPI; 2003-493399/46.
                                  WO2003045998-A2
                                                                                                                                                                                                                                                    Bourgeron T,
Gillberg C;
                                                                   05-JUN-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates ro a novel use of irinotecan or its derivative for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having agenome with a variant allele which comprises a multidaving resistance I (MDR1) polymuclectide. A composition of the invention has cytostatic activity. The present sequence is used in the exemplification of the invention.
                                                                                                                                                                                                               irinotecan; colorectal cancer; cervical cancer; gastric cancer;
lung cancer; ovarian cancer; pancreatic cancer; malignant glicma;
multidrug resistance 1; MDR1; cytostatic; ds; human; UGT1A1; MRP1; TOP1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New use of irinotecan for preparation of pharmaceutical compositions for treating cancer in subject having genome with variant allele comprising multidrug resistance 1 polynucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 21;
                                                                                                                                                                                Human UGT1A1 variant allele sequence fragment SEQ ID NO:2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 6 A; 9 C; 6 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 3.3%; Score 14.2; DB 1;
Best Local Similarity 84.2%; Pred. No. 2.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 41; 104pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EPID-) EPIDAUROS BIOTECHNOLOGIE AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         336 GACCAGGCCGGCTGCTCT 354
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21 Greeresecesecretrer 3
                                                                       ADB92135 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                       23-JUL-2001; 2001EP-00117608.
24-MAY-2002; 2002EP-00011710.
                                                                                                                                                                                                                                                                                                                                                                                                      23-JUL-2002; 2002WO-EP008220.
                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Heinrich G, Kerb R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-342400/32.
                                                                                                                                                                                                                                                                                                                              WO2003013535-A2.
                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                             04-DEC-2003
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                                                                                                          ADB92135
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Leboyer M;

Quach H, Betancur C,

Jamain S,

Homo sapiens

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                                                                                                                                                                                                                                                                                                                                              ss; gene; atherosclerotic lesion; antiatherosclerotic; cerebroprotective; antianginal; thrombolytic; cardiant; ophthalmological; neuroprotective; nephrotropic; vasotropic; atherosclerosis; stroke; angina; thrombosis; myocardial infarction; ischaemic heart disease; transplantation-induced sclerosis; intermittent claudication; diabetes; peripheral artery disease; congestive heart failure; retinopathy; neuropathy; thrombosis.
                                                                                                                                                                                                                                                                                                                DNA oligo (SeqID 93) encodes peptide that binds atherosclerotic lesions
                                                     Gaps
                                                     ö
              3.3%; Score 14.2; DB 1; Length 21; 84.2%; Pred. No. 2.7e+02; ive 0; Mismatches 3; Indels
                                                                                     41 AGATGGCCACCACTCAGAG 59
20 AGAAGGCCATCATTCAGAG 2
                                                                                                                                                                                                                BP.
Query Match
Best Local Similarity
                                                                                                                                                                                                                ADE77842 standard; DNA; 21
                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                  29-JAN-2004
                                                                                                                                                                                                                                                   ADE77842;
                                                                                                                                                                            RESULT 149
ADE77842/c
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Synthetic

nootropic; neuroleptic; tranquillizer; gene therapy; synaptogenesis; mutation; neurological disease; mental disorder; psychiatric illness; autism; Asperger syndrome; schizophrenia; autism; Asperger syndrome; schizophrenia; attention deficit hyperactivity disorder; ds; ss; primer.

Human HNL4X/Y gene PCR primer #4.

(first entry)

18-DEC-2003

ADC24720/ ID ADC2 AC ADC2 XX ADC2 XX ADC2 XX IB-D DE Huma XX NOOC XX MOOC X

ADC24720;

ADC24720 standard; DNA; 21 BP.

RESULT 148

WO2003014145-A2.

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This invention relates to novel isolated peptides that selectively bind to mammalian atherosclerotic lesions and as such can be used to detect to mammalian atherosclerotic lesions and as such can be used to detect to the in vivo identification of such peptides by using phage display the in vivo identification of such peptides by using phage display conditions of beptides. Display the peptides of perhotogical conditions of the conditions of the endothelial tissue occurs by administration of a peptide conjugated to reporter molecule or therapeutic agent. As such, these peptides can be careforted variously as antiatherosclerotic, cerebroprotective, antianginal, thrombolytic, cardiant, optibalmological, neuroprotective, oppides as useful for treating atherosclerosis, as well as identifying the location and severity of an atherosclerosis, as well as identifying the location and severity of an atherosclerosis, myocardial infarction, ischaemic heart disease, transplantation-induced sclerosis and confinence, it is associated with diabetes, which in turn can lead to peripheral artery disease, congestive heart failure, retinopathy, neuropathy, nephropathy or thrombosis. This colliponucleotide sequence, isolated from a combinatorial phage display collibrary, encodes a peptide that binds to atherosclerotic lesions, the aim
                                                                                                                                                                                                                                                                                                Novel peptide which selectively bind to mammalian atherosclerotic lesions, useful for treating atherosclerosis in a mammal, and for identifying location of atherosclerotic lesion in mammal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 6 A; 4 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                               Claim 16; SEQ ID NO 93; 286pp; English
                                                                                                                                                                                                             Liu C, Edgington TS, Prescott MF;
                                                                                                                                      (NOVS ) NOVARTIS AG.
(NOVS ) NOVARTIS PHARMA GMBH.
(SCRI ) SCRIPPS RES INST.
                                                                 09-AUG-2002; 2002WO-EP008942.
                                                                                                      10-AUG-2001; 2001US-0311507P
                                                                                                                                                                                                                                                WPI; 2003-278468/27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    of the invention.
                                                                                                                                                                                                                                                                     P-PSDB; ADE77843
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-JUL-1999
                                20-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAX64556;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         150
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Gaps
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Query Match 3.3%; Score 14.2; DB 1; Length 21; Best Local Similarity 84.2%; Pred. No. 2.7e+02; Matches 16; Conservative 0; Mismatches 3; Indels
                                                                                                                230 CAAATCGGGAGGCTGCTTC 248
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Human B7-1 hammerhead ribozyme target SEQ ID NO:1188. 21 CAAATCAGGAGTCTGATTC 3 AAX64556 standard; RNA; 15 BP. (first entry)

Arthritic condition; graft tolerance; immune response; target; cleavage; hammerhead ribozyme; hairpin ribozyme; human; rabbit; mouse; collagenase; stromelysin; synovial membrane; joint; arthritis; osteoarthritis; rheumatoid arthritis; autoimmune disease; allergy; inflammation; diagnosis; ss.

Homo sapiens

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Bhzymatic nucleic acid molecules having a hammer-head motif - used for the treatment of arthritis, induction of graft tolerance or treatment of
                                                                                                                                                              Stinchcomb DT, Jarvis T, Draper K, Pavco P;
Gustofson J, Usman N, Wincott F, Matulic-Adamic J;
Thompson JD, Modak A, Burgin A;
                                                                                                                                                                                                                                              Claim 10; Page 166; 307pp; English.
                                                   94US-00354920.
94US-00363254.
94US-003908324.
95US-00426124.
95US-00432874.
95US-000951P.
95US-000951P.
95US-000951P.
95US-000951P.
                                      95WO-US015516
                                                                                                                                                 (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                               auto-immune diseases.
                                                                                                                                                                                                WPI; 1996-300653/30.
                                                                                                                                                                Beigelman L,
Mcswiggen J,
Karpeisky A,
                                                   13-DEC-1994;
23-DEC-1994;
23-DEC-1995;
17-FEB-1995;
20-ARY-1995;
04-MAY-1995;
07-JUL-1995;
07-JUL-1995;
07-JUL-1995;
       WO9618736-A2
                                     22-NOV-1995;
                      20-JUN-1996.
                                                                                                                                  05-OCT-1995
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The present invention describes a novel enzymatic nucleic acid (ENA)

having a hammerhead motif (HM) comprising: (i) at least S ribose residues

(ii) a 2'-C-allyl modification at position 4 of the ENA; (iii) at least

(c ten 2'-O-methyl modifications; and (iv) a 3'-end modification. The ENA's

ccan inhibit collagenase and stromelysin production in the synovial

membrane of joints for the treatment or prevention of arthritis.

CC membrane of joints for the treatment or prevention of arthritis.

CC membrane of joints for the treatment or prevention of arthritis.

CC membrane of joints for the treatment or prevention of arthritis.

CC membrane of joints for the treatment or prevention of arthritis.

CC membrane of joints for the treatment of a donor to induce tolerance to a used to treat antigen presenting acels of a donor to induce tolerance of enhancing graft tolerance or for treating autoimmune disease, and for treating allergies and other inflammatory conditions. The ENA's can also be used in diagnosis. Ribozyme therapy impacts on the expression of stromelysin which accompany treatment with retinoids and dexamethasone.

CC The concentration of ribozyme required to affect a therapoutic treatment con specific. The present sequence is used in the exemplification of the present invention

Gaps ö Query Match
3.3%; Score 14; DB 1; Length 15;
Best Local Similarity 64.3%; Pred. No. 1.4e+02;
Matches 9; Conservative 5; Mismatches 0; Indels Sequence 15 BP; 2 A; 3 C; 5 G; 0 T; 5 U; 0 Other;

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HIV-1 Group O isolate HAM112 PCR primer env25R. AAX56095 standard; DNA; 18 BP. 401 GGTCTTCTACGTGA 414 2 GGUCUUCUACGUGA 15 15-JUL-1999 (first entry) AAX56095; RESULT 151 g RXXXXXXXX ઠે

New rapid assay for antibodies to HIV-1 groups O and M, and HIV-2 - can be used in field assay, requiring no electricity and less specialised equipment.

Example 2; Page 70; 104pp; English.

Hickman RK, Varitek V, Necklaws EC; Devare SG;

Vallari AS, Hackett JR, Golden AM, Brennan CA, WPI; 1999-190224/16.

(ABBO ) ABBOTT LAB. 15-AUG-1997;

97US-00912129

07-AUG-1998;

WO9909410-A2

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The present invention describes (A) an isolated HIV-1 Group O env polypeptide. Also described are: (1) an isolated HIV-1 Group O env polypeptide comprising an immunoreactive portion of a polypeptide as in (A) or (1); (2) a polymuclectide (PN) encoding a polypeptide as in (A) or (1); (3) an antigen construct comprising a first HIV-1 Group O env polypeptide (A) an antigen construct comprising a fusion of at least one HIV-1 Group O env polypeptide with at least one HIV-1 Group O env polypeptide with at least one HIV-1 Group O env polypeptide with at least one additional HIV-1 polypeptide; (5) an antigen construct comprising a fusion of a first HIV-2 env polypeptide; (6) an antigen construct of a first HIV-2 env polypeptide; (6) an antigen construct as in (3)-(6), (8) an expression vector comprising a mitigen construct as in (3)-(6), (8) an expression vector comprising a pN as in (7); (9) an immunoassay kit for the detection of antibodies to HIV-1 comprising an antigen construct as in (3)-(6). The antigen constructs can be used for the detection of anti-HIV-1 antibodies in test samples. They can also be used as immunogens to produce antibodies. The antibodies can be used to polypeptides, for therapy and for detection of HIV polypeptides, for therapy and for detection of HIV polypeptides.
                                                                                                                                                                                                                                                                                                                                                                                 New isolated HIV-1 Group O env polypeptides - used for the detection of anti-HIV antibodies and for the production of antibodies for use in detection, purification and therapy.
HIV; human immunodeficiency virus; antigen; detection; antibody; differentiation; Group O; env; immunogen; immunoassay; ss.
                                                                                                                                                                                                                                                                                                       Hackett JR, Yamaguchi J, Golden AM, Brennan CA, Hickman RK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ouery Match 3.3%; Score 14; DB 1; Length 18; Best Local Similarity 87.5%; Pred. No. 2.1e+02; Matches 14; Conservative 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 18 BP; 3 A; 7 C; 3 G; 4 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 2; Page 85; 138pp; English.
                                                       Synthetic.
Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              264 GTGCACCTGGAGCAGG 279
|:||||||||||||||||1|||1|||1||16 GYGCACCTGGAGTAGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAX37210 standard; DNA; 18 BP
                                                                                                                                                                                            98WO-US017014.
                                                                                                                                                                                                                               97US-00911824.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-JUL-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                              WPI; 1999-190167/16.
                                                                                                                                                                                                                                                                    (ABBO ) ABBOTT LAB.
                                                                                                                                                                                            17-AUG-1998;
                                                                                                                                                                                                                                 15-AUG-1997;
                                                                                                              WO9909179-A2
                                                                                                                                                     25-FEB-1999,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            셤
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The invention relates to a rapid assay for simultaneous detection and differentiation of antibodies to HIV-1 groups O and M, and HIV-2. The method comprises (a) contacting the sample with a strip containing at method comprises (a) contacting the sample with a strip containing at cleast one immobilised capture reagent par analyte and on which the sample conditions sufficient to form capture reagent par analyte complexes, and (b) determining the presence of analyte(s) by detecting a visible colour change at the capture reagent site on the strip wherein the capture comprises a polypeptide shown in AAV66983-84; and that for HIV-2 comprises the polypeptide shown in AAV66981. The invention is used to screen patients for antibodies to HIV-1 types O and M, and HIV-2. The invention will be particularly useful in the invention provides a screening method which is faster and requires comprises the polypeptide shown in the present patients for antibodies to HIV-1 types O and M, and HIV-2. The invention will be particularly useful in places and situation where equipment and/or electricity is not available. The invention provides a screening method which is faster and requires cless equipment than prior art methods. Sequences AAX37195-X37222 crepresent primers used for determining the env seqeunce of the HIV-1 course of group O isolate HAM112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; secreted protein; immunestimulant; immunesuppressant; virucide; antibacterial; antifungal; cytostatic; antinflammatory; dermatological; antidiabetic; antiartheumatic; antirheumatic; protozoacide; antidiabetic; antiartheumatic; protozoacide; antithyroid; immune deficiency; autorection dimmunedeficiency; SCID; infection; HIV; hepatitis; malaria; autorimmune disorder; systemic lupus; connective tissue disease; multiple sclerosis; erythematosis; rheumatoid arthritis; autorimmune pulmonary inflammation; asthma; quillain-Barre syndrome; autorimmune thyroiditis; myasthenia strais autorimmune thyroiditis; myasthenia agravis; autorimmune inflammatory eye disease; allergy; hybridisation; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3.3%; Score 14; DB 1; Length 18;
87.5%; Pred. No. 2.1e+02;
/ative 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human secreted protein clone ye90_1 probe SEQ ID NO:201.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 18 BP; 3 A; 7 C; 3 G; 4 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAA16738 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16-JUN-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity 87.5
Matches 14, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAA16738;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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SXCCCCCCCCCCCCXXXXIIIXXBXXIIXXBXXBXAXXBXXBXXBXXBXXBXXBXXBXXBX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ઠે
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HIV-1; HIV-2; immobilised capture reagent; capillary action; screening antibody; assay; env protein; PCR primer; ss.

Synthetic. Human immunodeficiency virus 1.

HIV-1 env sequence determining primer.

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New polynucleotides encoding secreted proteins, which may have e.g. nutritional, chemokine, immune stimulating or suppressing, hematopoiesis regulating, tissue growth, activin/inhibin antiinflammatory or tumor inhibition activity.
                                                                                                                                                  Collins-Racie LA, Evans C;
Steininger RJ, Spaulding V;
                                                                                                                                                                                                                                 Disclosure; Page 627; 641pp; English.
                                                                                                                                                K, Mccoy JM, Lavallie ER,
D, Treacy M, Agostino MJ,
, Clark HF, Fechtel K;
                                                                98US-0096815P.
98US-0099229P.
98US-0105368P.
99US-0119534P.
99US-0119531P.
99US-0120575P.
99US-013020P.
                                           99WO-US018298
                                                                                                                                  GEMY ) GENETICS INST INC.
                                                                                                                                                                                WPI; 2000-205979/18.
             WO200009552-A1.
Homo sapiens
                                           13-AUG-1999;
                                                                                                                     11-AUG-1999;
                            24-FEB-2000.
                                                                                                     18-FEB-1999
                                                                                 23-OCT-1998
                                                                                                                                                          Merberg D,
Wong GG,
                                                                                                                                                  Jacobs K,
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AAA16618 to AAA16697 encode the human secreted proteins given in AAY94898

to AAY94980, isolated from human adult brain, adult thyroid, adult
retina, foetal arcinoma, adult blood, adult heural, foetal kidney, adult
placenta, adult testis, whole embryo, adult cartilage, kidney, foetal
adult bladder, cDNA libraries. The polymicleotides and proteins are
predicted to have biological activities which would make them suitable
for treating, preventing or ameliorating medical conditions in humans and
animals. The polymicleotides can be used as markers for tissues in which
the protein is preferentially expressed, as molecular weight markers on
Southern gels, and as chromosome markers or tags to identify chromosomes
or to map gene positions. The proteins can be used in the treatment of
immunedeficiency (SCID), as well as viral, bacterial, fungal and other
infections. These infections include human immunodeficiency virus (HIV),
candidissis. The proteins can be used to treat autoimmune disorders such
as connective tissue disease, multiple sclerosis, systemic lupus
candidissis. The proteins can be used to treat autoimmune disorders activing activity inflammation,
dillain-Barre syndrome, autoimmune thyroiditis, insulin dependent
diabetes mellitus, myasthenia gravis, graft-versus-host-disease and
autoimmune inflammatory eye disease. The proteins can also be used to
treat allergic conditions, such asthma. AAA16688 to AAA16774 represent
probes for the human secreted proteins from the present invention

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0
                                 3.3%; Score 14; DB 1; Length 18; 00.0%; Pred. No. 2.1e+02;
                                                                 0; Indela
Sequence 18 BP; 5 A; 5 C; 6 G; 2 T; 0 U; 0 Other;
                                                                  0; Mismatches
                                                  100.0%;
                                                Local Similarity 100.
nes 14; Conservative
                                 Ouery Match
                                                                  Matches
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AAZ90302/c ID AAZ90302 standard; DNA; 18 BP. XX RESULT 154

(revised)
(first entry) 15-SEP-2003 22-MAY-2000 

HIV-1 env PCR primer env25R, SEQ ID NO:77.

HIV-1 group 0; env; gpl20; gp41; glycoprotein; monoclonal antibody; immunoassay; positive control; affinity purification; therapeutic; antigen; expression construct; PCR primer; ss.

Human immunodeficiency virus 1; group O isolate HAM112.

WO200004383-A2.

27-JAN-2000.

99WO-US015469. 09-JUL-1999;

98US-00115171 14-JUL-1998;

(ABBO ) ABBOTT LAB.

Scheffel JW, Hackett JR, Tyner JD, Hickman RK;

WPI; 2000-171290/15.

Novel monoclonal antibodies useful as positive control reagent for detecting human immunodeficiency virus infections and diagnosing, evaluating or prognosing viral disease.

Example 2; Page 37; 148pp; English.

The invention relates to anti-HIV-1 group O monoclonal antibodies, which may be used as positive control reagents in immunoassays to detect and differentiate HIV-1 infections. The invention also encompasses a monoclonal antibody which binds specifically to an HIV-1 group O antigen, which has no more than 15% cross reactivity to a corresponding antigen selected from HIV-1 group M antigens and HIV-2 antigens; and a method of using a monoclonal antibody as a positive control reagent in an immunoassay for the detection of anti HIV-1 group O antibodies. The monoclonal antibodies are useful as positive control reagents in an immunoassay involve coupling a monoclonal antibodies. Such immunoassays involve coupling a monoclonal antibodies. Such immunoassays involve coupling a monoclonal antibodies. Such cantibodies of the invention would be used to ensure that the reagents provided to detect HIV-1 group O antibody with HIV group-1 antibodies of the invention would be used to ensure that the reagents provided to detect HIV-1 group O antibody were performing properly. The monoclonal antibodies can be inmobilised on a marita and so constitutes or biological tissues. The monoclonal antibodies can also be used for generating chimeric antibodies for therapeutic use. Different synthetic, recombinant or purified antibodies for therapeutic use. Different epitopes of HIV antigens can be used in combination. The monoclonal antibodies are also useful for differentiating HIV-1 group O antigens of from HIV-2 monoclonal antibodies are also useful for differentiating HIV-1 group O antigens of generate and amplify obNA encoding the native env protein of HIV-1 group O, isolate HAM12. Sequences AA290301 represent PCR primers used in an exemplification of the present invention to generate expression constructs comprising HIV-1 group O envention.

C sequence AA290303 represents a primer of undefined function. (Updated on

Sequence 18 BP; 3 A; 7 C; 3 G; 4 T; 0 U; 1 Other;

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Gaps

Gaps ö 3.3%; Score 14; DB 1; Length 18; 87.5%; Pred. No. 2.1e+02; 1; Mismatches Query Match Best Local Similarity 87.59 warches 14; Conservative

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264 GTGCACCTGGAGCAGG 279

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16 GYGCACCTGGAGTAGG 1

rng.res

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8
Page
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Sequences AAC92738-C92817 represent antisense oligonucleotides targetted to the heterogeneous muclear ribonucleoprotein A1 (hnRNP A1) gene, which inhibit its expression. The antisense oligonucleotides were designed to target different regions of the human hnRNP A1 mRNA, and were analysed for their effect on hnRNP A1 mRNA levels by quantitative real-time PCR. hnRNP A1 (also known as heterogeneous nuclear ribonucleoprotein core protein A1 and processing (including alternative spliting) of newly transport and processing (including alternative spliting) of newly transport and processing (including alternative spliting) of newly transport and shutles continuously between the nucleus and the cytoplasm acting as an earrier protein for mRNAs. hnRNP A1 also participates in telomere biogenesis, with low levels of hnRNP correlating with shortened telomeres. In addition, hnRNP A1 has also been classified as an apoptosis associated protein on the basis that it is specifically cleaved into three fragments during antibody-mediated apoptosis. Due to its ability to control spliting events, particularly donor splice site selection, hnRNP control spliting events, particularly donor splice site selection, hnRNP control spliting events, particularly donor splice site selection, hnRNP control spliting events, particularly donor splice site selection, hnRNP control spliting events, particularly donor splice site selection, hnRNP control spliting events, particularly donor splice site selection, hnRNP control spliting events, particularly events on and treatment of the invention are useful for diagnosis, prevention and treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel antisense compound targeted to human hnRNP A1 which specifically hybridizes with and inhibits the expression of human hnRNP A1, useful for modulating the expression of hnRNP A1 in cells.
                                                    Human hnRNP A1; heterogeneous nuclear ribonucleoprotein A1; heterogeneous nuclear ribonucleoprotein core protein A1; p40CRS; mRNA processing; transport; stabilisation; alternative splicing; donor splice site selection; telomere biogenesis; oncogenesis; apoptosis-associated protein; cancer; tumour formation; expression inhibition; phosphorothioate; antisense oligonucleotide; ss.
                   Human hnRNP Al phosphorothioate antisense oligonucleotide, SEQ ID NO:57.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             conditions associated with hnRNP Al expression, such as cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
3.3%; Score 14; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 3 A; 8 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human gene single nucleotide polymorphism #2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 3; Col 41-42; 38pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         В.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAF97242 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19 TGGCGGTGGAGGCC 6
                                                                                                                                                                                                                                                                                                                                                                                               (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                       Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-090484/10.
                                                                                                                                                                                                                                                                                                                 27-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                       27-OCT-1999;
                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-JUN-2001
                                                                                                                                                                                                                                      US6165789-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                       Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAF97242;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ7401 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the pharmaceutical ascendical secting on a disease as well as other treatment. N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3056, 3157, 3227, 3297 and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel biallelic markers used to construct a high density disequilibrium
                                                                                                                                                                       Human biallelic marker downstream amplification primer SEQ ID NO:8409.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                        Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyping; hybridisation; identification; characterisation; amplification; single nucleotide polymorphism; SNP; PCR primer; diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 3 A; 5 C; 3 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chumakov I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 8; Page 2023; 2745pp; English.
                                                       AAZ74053 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98US-0082614P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          99WO-IB000822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98US-0109732P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TGAAAGCAGAGAAC 218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAC92785 standard; DNA; 20
                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cohen D, Blumenfeld M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   map of the human genome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-013267/01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21-APR-1998;
23-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          21-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                          WO9954500-A2
                                                                                                                                   10-SEP-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                  28-OCT-1999
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Best Local S:
Matches 14
                                                                                             AAZ74053;
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AAC92785/c
ID AAC9278
XX
AC AAC9278
XX
DC AAC9278
              RESULT 15
AAZ74053/
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Gaps

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pulmonary embolism; paternity test; ds

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The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclarosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thromboembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also cuseful in forenais, paternity testing, genetic analysis and phenotype correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
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               Human; variant thrombospondin 1; variant thrombospondin 4; SNP; polymorphism; vascular disease; coronary artery disease; forensics; myocardial infarction; atherosclerosis; stroke; venous thromboembolism; pulmonary embolism; paternity test; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bolk S, Daley GQ, Mccarthy JJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and atherosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 3.3%; Score 14; DB 1; Length 21; Best Local Similarity 100.0%; Pred. No. 3e+02; Matches 14; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 7 A; 4 C; 8 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gargill M, Ireland JS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example; Page 184; 242pp; English.
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AAF97748 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                                                                                                                  10-SEP-1999; 99US-0153357P.
26-JUL-2000; 2000US-0220947P.
16-AUG-2000; 2000US-025724P.
                                                                                                                                                                                                                                                                                                                 07-SEP-2000; 2000WO-US024503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20 GGTGACCGAGGGCT 33
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-226749/23.
                                                                                                                                                                                                                                          WO200118250-A2
                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                              15-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lander ES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAF97748;
                                                                                                                                                                  Variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 158
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The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various oblymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart disease, stroke, peripheral vacular diseases, venous thromboembolism pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Probe; quantification; human; GTP binding protein; G protein; alpha subunit; specific mRNA; detection; hybridisation; diagnosis; pathophysiology; disease state; hereditary; cancer; infectious; osteodystrophy; pituitary tumour; acromegaly; melanoma cells; diabetes; PCR; polymerase chain reaction; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                      Mccarthy JJ;
                                                                                                                                                                                                                                                                                                                                                                                                                  Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and atherosclerosis.
                                                                                                           /standard_name= "single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ..
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.3%; Score 14; DB 1; Length 21; 100.0%; Pred. No. 3e+02; ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                      S, Daley GQ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mouse D MUSJUNDA, MUSJUNDR/B-1258 jun-B specific probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 5 A; 4 C; 6 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                      Lander ES, Gargill M, Ireland JS, Bolk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AA047598 standard; cDNA to mRNA; 17 BP.
                                                                                                                                                                                                                                                                                                      (WHED ) WHITEHEAD INST BIOMEDICAL RES. (MILL-) MILLENNIUM PHARM INC.
                                                             Location/Qualiflers
replace(11,T)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example; Page 218; 242pp; English.
                                                                                                                                                                                                                                        10-SEP-1999; 99US-0153357P.
26-JUL-2000; 2000US-0220947P.
16-AUG-2000; 2000US-025724P.
                                                                                                                                                                                                             07-SEP-2000; 2000WO-US024503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          93 ATCACCACGTCTGA 106
|||||||||||||||||
19 ATCACCACGTCTGA 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 100..
                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-226749/23.
                                                                                                                                              WO200118250-A2
                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAR-2003
26-JAN-1994
                                                                                                                                                                            15-MAR-2001
                                                                 Key
Variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAQ47598;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 159
AAQ47598/c
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rng.res
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The sequences given in AAQ47594-603 show regions of homology between jun sequences and the jun-B specific probe B-1258 which may be of use as jun-B specific probe B-1258 which may be of use as jun-B specific probes. They were used in the method of the invention for the detection and quantification of markha in a sample without the need to purify the mRNA from cells. The claimed method comprises identifying a polynucleotide sequence unique to the mRNA, and immobilising an oligomer complementary to this sequence to an insoluble support. The sample is components are washed from the support such that the unique sequence will hybridise to the bound oligomer and bound RNA is labelled in such components are washed from the support and bound RNA is labelled in such a may that the label is inforced note the support relative to the amount of mRNA on the support. The amount of bound label is then amount of determined. This method can be used for the reliable, rapid, simultaneous determined. This method can be used for the reliable, rapid, simultaneous contention of multiple varieties of mRNA. It may be used for diagnosing and recognition of pathophysiology of various disease states, eg. hereditary diseases, cancer, and infectious diseases. Getiagnosing and recognition of pathophysiology of various disease (of thought to be involved in causing various diseases. Getiagnosing and recognitions of pathophysiology of various disease states. Geticiency of Ge proteins is the molecular basis of hereditary contents multiple varieties of proteins are also involved in invasive and metastatic melanoma cells, and diabetes. See also AAQ47381-666. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Quantitating messenger RNA in sample - using immobilised-polynucleotide having sequence complementary to sequence unique to the MRNA.
                                                                                                                                                                                                                                                                                                                                                      Akitaya T, Cooper A, Mitsuhashi M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 9; Page 71; 177pp; English.
                                                                                                                                                                                                                                                                                    (HITB ) HITACHI CHEM CO LTD.
(HITB ) HITACHI CHEM RES CENT INC.
                                                                                                                                                                                     92US-00827208.
92US-00857059.
92US-00974409.
                                                                                                                                     93WO-US000977
                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1993-258695/32.
                                                                                                                                                                                     29-JAN-1992;
24-MAR-1992;
12-NOV-1992;
                                           WO9315221-A1
                                                                                                                                          29-JAN-1993;
                                                                                         05-AUG-1993
Synthetic.
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## Sequence 17 BP; 2 A; 8 C; 5 G; 2 T; 0 U; 0 Other;

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Gaps
                                                              ;
0
Ouery Match
3.2%; Score 13.8; DB 1; Length 17;
Best Local Similarity 88.2%; Pred. No. 2.1e+02;
Matches 15; Conservative 0; Mismatches 2; Indels
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AAZ39286 standard; DNA; 17 BP. RESULT 160

11-FEB-2000 (first entry) AAZ39286;

Probe for typing HLA allele B*1406.

Human leukocyte antigen; HLA; allele; HLA-B*3913; HLA-B*1406; human; HLA-B*51; HLA-DRB1*0820; HLA-DRB1*04; HLA-DRB4*01; allele typing; exon; major histocompatibility complex; MHC; probe; 98. SXXXXXXXXXXXXXX

Synthetic

The invention provides polymucleotides corresponding to exon 2 and exon 3 of human leukocyte antigen (HLA) alleles HLA-B*3913, HLA-B*1406 and HLA-B*51 and exon 2 of HLA alleles HLA-DEB1*0820, HLA-DEB1*04 and HLA-DEB1*01. The polymucleotides are useful for typing the above HLA alleles in a sample, especially by a method that comprises (a) amplifying all/part of the relevant sequence using at least one primer pair, and (b) hybridizing the amplified product to set of probes specifically hybridizing to target regions comprising one or more polymorphic nucleotides of the sequence, to determine the absence or presence of the anception preferred primer and/or at least one preferred primer and/or at least one preferred primer and/or at least one preferred probe and (b) for detecting the protein fragment encoded by the probe and (b) for detecting the protein fragment encoded by the pinding specifically to the protein fragment are provided. The polymucleotides also enable the isolation of the complete respective genes from a human genomic library New polynucleotides for human leukocyte antigen, HLA, allele fragments, useful for typing HLA alleles. Query Match 3.2%; Score 13.8; DB 1; Length 17; Best Local Similarity 88.2%; Pred. No. 2.16+02; Matches 15; Conservative 0; Mismatches 2; Indels Sequence 17 BP; 3 A; 5 C; 6 G; 3 T; 0 U; 0 Other; Rossau R; Claim 16; Page 19; 62pp; English. 98BP-00870088 (INNO-) INNOGENETICS NV De Canck I, Mersch G, WPI; 1999-634008/54. 20-APR-1998; 

298 AGGACCTGAGCCCCGGG 314 AGGACCIGAGCICCIGG 17 g

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Ribozyme, erythropoietin, granulocyte colony stimulating factor; interferon alpha, ss.

Hammerhead ribozyme substrate #3478.

WO9954496-A2 Homo sapiens

28-OCT-1999

99WO-EP002614 19-APR-1999;

Mcswiggen J;

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The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 orphan receptor, EAR3/CHD-TR-1, the GATA transcription factor gene, IRF-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoietin, granulocyte colony stimulating factor protein and interferon alpha
                                                                                                                                                            Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17 BP; 2 A; 10 C; 3 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                          Claim 54; Page 136; 164pp; English.
Zwick M, Pavco P,
                                                                              WPI; 2000-647423/62.
ij
Blatt
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Score 13.8; DB 1; Length 17;
Pred. No. 2.1e+02;
0; Mismatches 2; Indels
                                                              35
         3.2%;
                                                                                      17 GGGGGACCGAGGGCTTG 1
                                                              19 GGGTGACCGAGGGCTGG
Query Match
Best Local Similarity 88.2.
Best Local Similarity
Local Similarity
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Gaps ö

> ABK00841 standard; RNA; 17 BP (first entry) Human NOGO Inozyme #111. 12-MAR-2002 ABK00841; RESULT 1

Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; mootropic; neuroprotective; antiparkinsonian; muscular; D20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; DNAzyme; inozyme; G-cleaver; amberzyme; inizyme; I-ymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphomy; leukaemia; human immunodeficiency virus; HIV associated NHL; mantle-cell lymphoma; MCI; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; Parkinson's disease; ataxia; Huntingcon's disease; creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.

09-FEB-2001; 2001WO-US004273 11-FEB-2000; 2000US-0181797P. 28-FEB-2000; 2000US-0185516P. 06-MAR-2000; 2000US-0187128P. WO200159103-A2. sapiens. 16-AUG-2001. Synthetic.

RIBOZYME PHARM INC. BLATT L. MCSWIGGEN J. CHOWRIRA B M. (RIBO-) I (BLAT/) I (MCSW/) P

Chowrira BM; Mcswiggen J, ŭ Blatt

CHOM/)

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The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NGCD). The regulates expression of a neurite growth inhibitor gene (NGCD). The collate and processing and with a NGW motif) a d-cleaver (cleaving RNA with a NWN motif) processing an NCH motif), a d-cleaver (cleaving RNA with a NWN motif) processing an NCH motif), a d-cleaver (cleaving RNA with a NWN motif) processing an NCH motif), a d-cleaver (cleaving RNA with a NWN motif) processing an NCH motif), a d-cleaver (cleaving RNA with a NWN motif) processing an NCH motif) and cleaving RNA with a NWN motif) processing an NCH motify and inhomation and condition associated with the level of CD20 in the presence of a divalent cation that is preferably Mg^2+.

C Furthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of cleaving LNAMPhoma, leukaemia, HIV (human immunodeficiency virus) associated with, lymphocytic lymphoma, leukaemia, HIV (human immunodeficiency virus) associated NHL, mantle-cell lymphoma (NCL), immunocytema (NML), small B-cell lymphocytic lymphoma (NCL), immunocytema (NMC), small B-cell lymphocytic lymphoma (NCC), immunocytena (NCC), small active associated with, mantle-cell cargetting nucleic acid may be contacted with a cell to reduce NOGO gene in the cargetting nucleic acid may be contacted with a cell to reduce NOGO gene in the cell and treat a patient having a condition associated with the level of NOGO. The treatment may further comprise the use of one or more contact central nervous system (CNS) injury and cerebrovascular accident (CNA, stroke), Alzabeimer's disease, dementia, multiple sclerosis (NS), charapies. In particular, which comprise the use of one or more contact central nervous system (NS) injury and cerebrovascular accident central adisease, disease, demential metals adisease, disease, disease, demential cent
                                                                            Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sequence is an inozyme of the invention
                                                                                                                                                                                                                                  Claim 88; Page 79; 200pp; English.
                                                                                                                                                                         central nervous system injury.
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Gaps ô Query Match 3.2%; Score 13.8; DB 1; Length 17; Best Local Similarity 88.2%; Pred. No. 2.1e+02; Matches 15; Conservative 0; Mismatches 2; Indels 302 CCTGAGCCCCGGGGACC 318

Sequence 17 BP; 1 A; 9 C; 7.G; 0 T; 0 U; 0 Other;

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1 ccedcecccccedeacc 17 ABN05998 standard; DNA; 17 BP

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Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss. Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:5990.

(first entry)

29-MAY-2002

ABN05998;

Homo sapiens 

WO200192524-A2

06-DEC-2001

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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to hGDMLP-1 protein substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22.

The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence data for this patent did not form part of the printed of the sequence data for this patent did not form part of the printed of the printed of the present directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                         Hanzel DK, Rank DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ftp.wipo.int/pub/published_pct_sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; SEQ ID NO 5990; 214pp; English.
                                                                                                                                                                  30-JAN-2001; 2001MO-US000663.
30-JAN-2001; 2001MO-US000664.
30-JAN-2001; 2001MO-US000665.
30-JAN-2001; 2001MO-US000666.
30-JAN-2001; 2001MO-US000668.
30-JAN-2001; 2001MO-US000669.
30-JAN-2001; 2001MO-US000669.
30-JAN-2001; 2001MO-US000669.
                                                                                                      2000GB-00024263
2001WO-US000661
2001WO-US000662
25-MAY-2001; 2001WO-US016981
                                                                                      2000US-0236359P
                                                                                                                                                                                                                                                                                                                                                                                                                                       Gu Y, Ji Y, Penn SG,
                                                                                                                                                                                                                                                                                                                                                                                             (AEOM-) AEOMICA INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-179446/23.
                                                                                                        04-OCT-2000; 2
30-JAN-2001; 3
30-JAN-2001;
                                         26-MAY-2000;
21-SEP-2000;
27-SEP-2000;
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ö Gaps .. 0 / Match 3.2%; Score 13.8; DB 1; Length 17; Local Similarity 88.2%; Pred. No. 2.1e+02; res 15; Conservative 0; Mismatches 2; Indels Sequence 17 BP; 6 A; 2 C; 6 G; 3 T; 0 U; 0 Other; Query Match

352 TCTACAGCGACTTCCTC 368 17 TCTACATGGACTTCCTC 1 g ð

ABN07568 standard; DNA; 17 BP ABN07568; ABN07568
1D ABN0
XX
AC ABN0
XX
DT 29-M
XX
DE Huma

RESULT 164

29-MAY-2002 (first entry)

Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:7560.

New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1. Human; genome-derived myosin-like protein 1; GDVLP-1; hGDVLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss. Shannon ME; Chen W, Hanzel DK, Rank DR, 04-0CT-2000; 2000GB-00024263.
30-JAN-2001; 2001W0-US000661.
30-JAN-2001; 2001W0-US000662.
30-JAN-2001; 2001W0-US000663.
30-JAN-2001; 2001W0-US000666.
30-JAN-2001; 2001W0-US000666.
30-JAN-2001; 2001W0-US000666.
30-JAN-2001; 2001W0-US000669.
30-JAN-2001; 2001W0-US000669.
30-JAN-2001; 2001W0-US000669.
30-JAN-2001; 2001W0-US000669. 25-MAY-2001; 2001WO-US016981 2000US-0236359P Gu Y, Ji Y, Penn SG, WPI; 2002-179446/23. (AEOM-) AEOMICA INC. WO200192524-A2 21-SEP-2000; 27-SEP-2000; Homo sapiens. 26-MAY-2000; 06-DEC-2001 

Shannon ME;

Chen W,

Disclosure; SEQ ID NO 7560; 214pp; English.

The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify concleic acids can be used as probes to detect, characterise and quantify concleic acids can be used as probes to detect, characterise and quantify convide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP concentration and/or amount specifically of hGDMLP proteins, as specific blomolecule capture probes for surface-enhanced laser desorption ionisation, as therapout engagement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The concentration and skeleral muscle disorders. hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart of hGDMLP-1 sequence in the spenential cate for this patent did not form part of the printed capture data for this patent did not form part of the printed sequence data for this patent did not form part of the printed sequence associated with respective conformed the respective of a fitp.wipo.int/pub/published_pot_sequence

Sequence 17 BP; 6 A; 3 C; 7 G; 1 T; 0 U; 0 Other;

Gaps . 0 Query Match
3.2%; Score 13.8; DB 1; Length 17;
Best Local Similarity 88.2%; Pred. No. 2.1e+02;
Matches 15; Conservative 0; Mismatches 2; Indels

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Sequence 17 BP; 5 A; 2 C; 6 G; 4 T; 0 U; 0 Other;

at ftp.wipo.int/pub/published_pct_sequence

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Human, genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
                                                                                                                                                                                                                                                                                                              Shannon ME;
                                               Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:5989.
                                                                                                                                                                                                                                                                                                              Chen W,
                                                                                                                                                                                                                                                                                                              Hanzel DK, Rank DR,
                                                                                                                                                                                                    30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
ABN05997 standard; DNA; 17 BP
                                                                                                                                                                                                                                                     2001WO-US000668.
2001WO-US000669.
2001WO-US000670.
                                                                                                                                                                       2000US-0234687P.
2000US-0236359P.
                                                                                                                                                                                      2000GB-00024263
                                                                                                                                                                                                                                                                              2001US-0266860P
                                                                                                                                                25-MAY-2001; 2001WO-US016981
                                29-MAY-2002 (first entry)
                                                                                                                                                                                                                                                                                                              Gu Y, Ji Y, Penn SG,
                                                                                                                                                                                                                                                                                                                            WPI; 2002-179446/23.
                                                                                                                                                                                                                                                                                              (AEOM-) AEOMICA INC.
                                                                                                               WO200192524-A2.
                                                                                                                                                                                                                                                              30-JAN-2001;
30-JAN-2001;
05-FEB-2001;
                                                                                                Homo sapiens
                                                                                                                                                                       21-SEP-2000;
27-SEP-2000;
                                                                                                                                                                26-MAY-2000;
                                                                                                                                                                                               30-JAN-2001:
                                                                                                                                06-DEC-2001.
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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 uncleic acids in samples, as amplification substrates, to hGDMLP-1 nucleic acids in samples, as amplification substrates, to hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule and/or amount specifically proteins, as specific deficiency in hGDMLP-1 proteins, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromesome 22.

The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from MIPO New polypeptide, for raising antibodies that recognize hGDWLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDWLP-1. Disclosure; SEQ ID NO 5989; 214pp; English.

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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific blomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                                  Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
                                Gaps
                                                                                                                                                                                                                                                                       Human GDWLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:7562.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Shannon ME;
                                ö
3.2%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 2.16+02; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chen W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rank DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 7562; 214pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WG-US000670.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-OCT-2000; 2000GB-00024263.
30-JAN-2001; 2001MC-US000661.
30-JAN-2001; 2001MC-US000662.
30-JAN-2001; 2001MC-US000663.
30-JAN-2001; 2001MC-US000663.
                                                                353 CTACAGCGACTTCCTCA 369
                                                                                                                                                                             ABN07570 standard; DNA; 17 BP
                                                                                   17 CTACATGGACTTCCTCA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2000US-0234687P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAY-2001; 2001WO-US016981
                                                                                                                                                                                                                                            29-MAY-2002 (first entry)
                 Local Similarity 88.2 les 15; Conservative
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27-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
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                                                                                                                                                                                                             ABN07570;
   Query Match
                                                                                                                                                RESULT 166
                                Matches
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Gaps

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New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser descrption ionization, comprises human myosin-like protein hGDMLP-1.
and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as preheargeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polymuclectide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMMLP-1 sequence in the exemplification of the present invention. N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in abetronic format directly from WIPO at fitp.wipo.int/published_pot_sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:5991.
                                                                                                                                                                                                                                                                      Query Match

3.2%; Score 13.8; DB 1; Length 17;

Best Local Similarity 88.2%; Pred. No. 2.1e+02;

Matches 15; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                      Sequence 17 BP; 6 A; 3 C; 7 G; 1 T; 0 U; 0 Other;
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21-SEP-2000; 2000US-0234687P.
27-SEP-2000; 2000US-0236359P.
04-OCT-2000; 2000US-00264263.
30-JAN-2001; 2001WO-US000662.
30-JAN-2001; 2001WO-US0006663.
30-JAN-2001; 2001WO-US0006663.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000669.
                                                                                                                                                                                                                                                                                                                                                 387 GACGCCCCAAGAAGGT 403
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ABN05999 standard; DNA; 17 BP.
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Shannon ME;

Disclosure; SEQ ID NO 5991; 214pp; English

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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polymucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify protein variants having desired phenotypic improvements and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as specifically recognise hGDMLP-1 proteins, as specifically of hGDMLP proteins, as specific biomolecule and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as the response or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart of selection miscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at the present directly from WIPO at the present directly from WIPO
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3.2%; Score 13.8; DB 1; Length 17;
Best Local Similarity 88.2%; Pred. No. 2.1e+02;
Matches 15; Conservative 0; Mismatches 2; Indels
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30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
23-MAY-2001; 2001US-00864761.
09-OCT-2001; 2001US-0327898P.
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ABV79108
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wed Apr 21 12:35:21 2004
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The present invention relates to human testis expressed Patched like protein (HTPL, see ABV78759 to ABV78762 and ABB98519 to ABB98520). HTPL has two isoforms, with a few single base pair differences between the two. One of the single base pair changes introduces a premature stop codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL shared structural features extrongly imply that HTPL plays a role similar to that of Patched, and is a potential tumour suppressor. HTPL gene was important in regulating male germ cell development, and the HTPL gene was important in regulating male germ cell development, and the HTPL gene was mapped to human chromosome 10p12.1. HTPL and its coding sequence are useful for diagnosing a disorder caused by mutation in HTPL, and in therapy and manufacture of a medicament for treatment or prevention of such disorder associated with decreased expression or activity of human HTPL. Such disorders include disorders of testis, or adrenal, adult and foctal liver, bone marrow, brain, kidney, lung, placenta, prostate, skeletal muscle or colon function. HTPL proteins and nucleic acids are clinically useful diagnostic markers and potenial therapeutic agents for male infertility and cancer. The present oligonucleotide was used in an example from the invention Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL. Example 2; Page 110; 718pp; English. 

Sequence 17 BP; 1 A; 8 C; 6 G; 2 T; 0 U; 0 Other;

0; Gaps Query Match 3.2%; Score 13.8; DB 1; Length 17; Best Local Similarity 88.2%; Pred. No. 2.1e+02; Matches 15; Conservative 0; Mismatches 2; Indels

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ABV91035 standard; DNA; 17 BP ABV91035; ABV91035

23-DEC-2002 (first entry)

Human POSHL1 scanning oligonucleotide SEQ ID NO 1748.

Human; POSHL 1; SH3 domain; POSH-like signalling protein 1; oncogene; Rho GTPase; signal transduction; gene expression; cancer; vaccine; gene therapy; transgenic; ss. 

Homo sapiens

EP1239051-A2.

11-SEP-2002.

28-JAN-2002; 2002EP-00001165

30-JAN-2001, 2001WO-US000663. 30-JAN-2001; 2001WO-US000664. 30-JAN-2001; 2001WO-US000665. 30-JAN-2001; 2001WO-US000665. 30-JAN-2001; 2011WO-US000667. 30-JAN-2001; 2011WO-US000668. 23-MAY-2001; 2001US-00864761, 10-OCT-2001; 2001US-0328205P.

(AEOM-) AEOMICA INC.

WPI; 2002-684061/74

Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1.

The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino acids (SL, ABB83999), a sequence having 65% sequence identity to (S1), (S1), (S1), having 95% deviations, especially conservative substitutions or a fragment of the sequence comprising at least 8 contiguous amino acids. Human POSHL 11s a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPsses as well as downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (I) and nucleic acids (II) for identifying a specific binding partner. (I) and nucleic acids (II) conced by altered expression of human POSHL1 including diagnosing and treating cancer, they useful in the development of vaccines and (II) is treating cancer, they useful in the development of vaccines and (II) is useful for measuring and for surveying gene expression and creating treansgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples of the invention. Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to between by the European Patent Office 

Sequence 17 BP; 3 A; 7 C; 6 G; 1 T; 0 U; 0 Other;

Gaps ó 3.2%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 2.1e+02; vative 0; Mismatches 2; Indels

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338 CCAGGGCCGGCTGCTCT 354 17 CCAGGGCCGGCTGTGCT 1 ઠ 셤

ABL31539 standard; DNA; 17 BP. ABL31539;

21-MAR-2002 (first entry)

Human HLA genotyping oligonucleotide SEQ ID NO 1028.

Human, human leukocyte antigen, HLA, genotype, polymorphism, immunogenetic, transplantation, genetic disease, ss.

Homo sapiens

WO200192572-A1.

06-DEC-2001.

01-JUN-2001; 2001WO-JP004662.

01-JUN-2000; 2000JP-00164798.

(NISN ) NISSHINBO IND INC. (SYST-) SYSTEM RES INC.

Moriya S, Nishida M; Matsumura Y, Inoko H, Kagiya T, Ichihara T, WPI; 2002-122074/16. Human leukocyte antigen (HLA) typing, useful for judging HLA genotypes of individuals e.g. by determining immunogenetic differences when transplanting between them.

Example 2; SEQ ID NO 1748; 60pp + Sequence Listing; English.

Query Match Best Local Similarity 88.2%

RESULT 170 ABL31539

THE SECRET SECRE

The invention relates to a typing kit for judging human leukocyte antigen (RIA) genotype of a sample by hybridising a substrate on which 10-24 base oligomicolotides (AB103612-AB131809) originating in the sequences of genes e.g. belonging to HiA class I antigens on human genome and containing gene polymorphisms as alloantigens have been immobilised as primers for amplification of cleaved nucleic acids relating to gene polymorphisms. The method is useful for judging HiA genotypes of individuals by determining immunogenetic differences before transplanting between them, providing genetic information to decide compatibility of organ and tissue for transplantation e.g. of bone marrow, kidney, liver, pancreas, Langerhans islet in pancreas and cornea, susceptibility diagnosis of genetic diseases and identifying individuals Sequence 17 BP; 3 A; 5 C; 6 G; 3 T; 0 U; 0 Other; 

Claim 10; Page 288; 345pp; Japanese.

3.2%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 2.16+02; tive 0; Mismatches 2; Indels Query Match

298 AGGACCTGAGCCCCGGG 314 Local Similarity 88.2 es 15, Conservative

1 Accaccicacciccics 17

RESULT 171

ABL31778 standard; DNA; 17 BP (first entry) 21-MAR-2002 ABL31778; ABL31778 

Human HLA genotyping oligonucleotide SEQ ID NO 1267.

Human; human leukocyte antigen; HLA; genotype; polymorphism; immunogenetic; transplantation; genetic disease; ss.

Homo sapiens.

WO200192572-A1.

06-DEC-2001,

01-JUN-2001; 2001WO-JP004662.

01-JUN-2000; 2000JP-00164798.

NISN ) NISSHINBO IND INC.

(SYST-) SYSTEM RES INC.

Ichihara T, Matsumura Y, Moriya S, Nishida M; Kagiya T, Inoko H,

Human leukocyte antigen (HLA) typing, useful for judging HLA genotypes of individuals e.g. by determining immunogenetic differences when transplanting between them. WPI; 2002-122074/16.

Claim 10; Page 333; 345pp; Japanese.

The invention relates to a typing kit for judging human leukocyte antigen (HLA) genotype of a sample by hybridising a substrate on which 10-24 base oligonucleotides (ABL30512-ABL31809) originating in the sequences of genes e.g. belonging to HLA class I antigens on human genome and containing gene polymorphisms as alloantigens have been immobilised as primers for amplification of clasved nucleic acids relating to gene polymorphisms. The method is useful for judging HLA genotypes of individuals by determining immunopenetic differences before transplanting between them, providing genetic information to decide compatibility of organ and tissue for transplantation e.g. of bone marrow, kidney, liver,

pancreas, Langerhans islet in pancreas and cornea, susceptibility diagnosis of genetic diseases and identifying individuals ន្តដូន

Sequence 17 BP; 3 A; 5 C; 6 G; 3 T; 0 U; 0 Other;

Gaps ö Length 17; 2; Indels 3.2%; Score 13.8; DB 1; 88.2%; Pred. No. 2.1e+02; tive 0; Mismatches 2; 15, Conservative Query Match Best Local Similarity Matches

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298 AGGACCTGAGCCCCGGG 314 ò

1 AGGACCTGAGCTCCTGG 17

RESULT 172 ACA0777

7711/c ACA07771 standard; RNA; 17 BP.

ACA07771;

03-JUN-2003 (first entry)

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0; Gaps

NFKB sub-unit modulating zinzyme substrate #170.

Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; d-claaver; amberzyme; cancer; REL-A activity; breast cancer; human; oesophageal cancer; stomach cancer; bladder cancer; prostate cancer; bladder cancer; parcreatic cancer; oesophageal cancer; stomach cancer; bladder cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemocherapy; paclitexel; docetaxel; cisplatin; methotrexate; chemocherapy; paclitexel; docetaxel; cisplatin; methotrexate; cyclophosphanids; docrubin; fluorouracil cancer; REL-A-specific inhibitor; cyclophosphanids; cacrubin; fluorouracil carbophatin; edatrexate; remematoid arthritis; restenosis; Crohn's disease; obesity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; schaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; schaemia; transplant/graft rejection; reperfusion injury; glomerulonesphritis; allergic airway inflammation; inflammatory bowel disease; infection; se

Homo sapiens.

US2002177568-A1.

28-NOV-2002.

23-MAY-2001; 2001US-00864785

92US-00987132. 94US-00245466. 94US-00291932. 96US-00777916. 18-MAY-1994; 15-AUG-1994; 23-DEC-1996; 07-DEC-1992; 

(STIN/) STINCHCOMB D T. (MCSW/) MCSWIGGEN J (DRAP/) DRAPER K G.

Stinchcomb DT, Mcswiggen J, Draper KG;

WPI; 2003-340953/32.

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.

Claim 3; Page 40; 72pp; English.

The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor regulates expression of a sequence encoding a subunit of nuclear factor configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for tracting a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially Mg^2^+. The enzymatic and

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prostate, coloractal brain, oesophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paciltaxel, docetaxel, cipplatin, methotrexate, gemcitabine or radiation, fluorouracil carboplatin, methotrexate, gemcitabine or radiation therapy. The enzymatic and antisense nucleic cid molecules are also useful for treating inflammatory disease such as rheumatoid arthritis, restenosis, asthma, crohn's disease, diabetes, obsestiv, autoimmune disease, lupus, multiple sclerosis, transplant/graft rejection, gene therapy applications, ischaemia/reperfusion injury (central nervous system (CNS) and myocardial), glomeruhonephritis, september alway inflammatory bowel disease or infection. This sequence represents the substrate of a novel enzymatic nucleic acid molecule
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88.2%; Pred. No. 2.18+02;
iive 0; Mismatches 2; Indels
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Best Local 9
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Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ1; MDZ1; chromosome 7g22.1; chromosome 6p21.3-22.2; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15g26.1; cancer; developmental disorder; ss. New zinc finger-containing proteins and nucleic acids, useful in manifacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MD24, MD27 or MD212, e.g. cancer. Example 8; SEQ ID NO 400; 103pp; English. 30-JUL-2002; 2002EP-00016874. 02-AUG-2001; 2001US-00922181 Gu Y, Nguyen C; WPI; 2003-423107/40. (AEOM-) AEOMICA INC. Homo sapiens. EP1281758-A2. 05-FEB-2003. Shannon M, 

Claim 1; Page 298; 387pp; English

The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ1, MDZ12. MDZ3 is encoded at chromosome 7q22.1, MDZ4 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 15p11.2 and MDZ12 is encoded at chromosome 15p26.1, The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3,

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MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as theraputic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nucleic acid molecule, Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synchesis; antisense; enzymatic nucleic acid; hammerhead ribozyme; DNAzyme; inozyme; zinzyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; viral replication; degemerative; disease state; HBV infection; HCV infection; dirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                               Query Match 3.2%; Score 13.8; DB 1; Length 17; Best Local Similarity 88.2%; Pred. No. 2.1e+02; Matches 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                              HCV minus strand DNAzyme substrate sequence #1332.
                                                                                                                                                Sequence 17 BP; 3 A; 6 C; 2 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                               361 ACTICCICACTITCCTG 377
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                                                                                                                                                                                                                                                                                                                                                                  ACD63973 standard; RNA; 17 BP.
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08-UUN-2001; 2001US-00877478.
08-UUN-2001; 2001US-0296876P.
24-OCT-2001; 2001US-0335059P.
05-DEC-2001; 2001US-0337055P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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Roberts E;
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MACEJAK D.
MCSWIGGEN J.
MORKISSEY D.
PAVCO P.
LEE P.
DRAPER K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-229207/22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (BLAT) BLATT L.
(MACE) MACEJAK D.
(MCSW) MACEJAK D.
(MORK) MORRISSEY D.
(MORK) MORRISSEY D.
(PAVC) PAVO P.
(LEEP) LEE P.
(LEEP) LEE F.
(CRAP,) DRAPER K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200281494-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                 30-SEP-2003
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Draper K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           infection.
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                                                                                                                                                                                                                                                                                                                                 RESULT 174
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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HTV) or the synthesis, expression and/or stability of Hepatitis C virus (HTV) or the patitis B virus (HTV) RNA. The nucleic acid and such as hammerhead ribozymes, DNAzymes, and enzymatic nucleic acid such as hammerhead ribozymes. DNAzymes, inozymes, zinzymes, amberzymes, and G-cleaver ribozymes. DNAzymes, or are nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptase primer sequences, as well transcriptase and/or HBV reverse transcriptase primer sequences, as well on Na. The nucleic acids may be used to modulate the expression of HBV or genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that methods of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene expression such as cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence sequences disclosed in the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      invention
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Gaps ; 3.2%; Score 13.8; DB 1; Length 17; 82.4%; Pred. No. 2.1e+02; ive 1; Mismatches 2; Indels Sequence 17 BP, 5 A, 3 C, 8 G, 0 T, 1 U, 0 Other; 14; Conservative lest Local Similarity Matches

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AAZ39244

AAZ39244 standard; DNA; 18 AAZ39244; 

BP.

Probe for typing HLA allele B*3913.

(first entry)

11-FEB-2000

Human leukocyte antigen; HLA; allele; HLA-B*3913; HLA-B*1406; human; HLA-B*51; HLA-DRB1*0820; HLA-DRB1*04; HLA-DRB4*01; allele typing; exon; major histocompatibility complex; MHC; probe; 85.

Homo sapiens. Synthetic.

WO9954496-A2

28-OCT-1999.

99WO-EP002614. 19-APR-1999;

9BEP-00870088.

20-APR-1998;

(INNO-) INNOGENETICS NV.

De Canck I, Mersch G,

New polynucleotides for human leukocyte antigen, HLA, allele fragments, useful for typing HLA alleles. WPI; 1999-634008/54.

The invention provides polynucleotides corresponding to exon 2 and exon 3 of human leukocyte antigen (HLA) alleles HLA-B*3913, HLA-B*1406 and HLA-B*51 and exon 2 of HLA alleles HLA-DRB1*0820, HLA-DRB1*064 and HLA-DRB4*01. The polynucleotides are useful for typing the above HLA alleles in a sample, especially by a method that comprises (a) amplifying all/part of the relevant sequence using at least one primer pair; and (b) Claim 16; Page 18; 62pp; English.

hybridizing the amplified product to a set of probes specifically hybridizing to target regions comprising one or more polymorphic nucleotides of the sequence, to determine the absence or presence of the allele in the sample. Diagnostic kits for (a) typing the alleles comprising at least one preferred primer and/or at least one preferred primer and/or at least one preferred probe and (b) for detecting the protein fragment encoded by the polymucleotides, comprising an antiserum or ligand (e.g. antibody) binding specifically to the protein fragment are provided. The polymucleotides also enable the isolation of the complete respective genes from a human genomic library 8×36666666666888

Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

ö Gaps . 3.2%; Score 13.8; DB 1; Length 18; 88.2%; Pred. No. 2.3e+02; ive 0; Mismatches 2; Indels Query Match
Best Local Similarity 88.2'
Matches 15; Conservative

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AAZ35254 standard; DNA; 18

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(first entry) 27-MAR-2000 AAZ35254; 

Plant retroelement primer binding site version 2.

Retroelement, retrovirus, transgenic plant, gene transfer, primer binding site, soybean, ss.

Glycine max.

WO9960842-A2

02-DEC-1999.

99WO-US011858 28-MAY-1999; 98US-0087125P. 99US-00322478. 29-MAY-1998; 28-MAY-1999;

(WRIG/) WRIGHT (VOYTAS

Wright DA, Voytas DF; WPI; 2000-105586/09. New nucleic acid molecules for imparting agronomically significant characters to plants, especially soybean.

Claim 1(a); Page 72; 118pp; English.

This oligonucleotide represents a soybean retroelement primer binding site (version 2). The invention provides molecular tools in the form of site (version 2). The invention provides molecular tools in the form of retroelements and plants.

Methods are provided for introducing the retroelements into cells, especially when the retroelement carries at least 1 agronomically, capacitation characteristic. In a preferred method, a helper cell line which expresses gag, pol and env sequences is used to enable transfer of which expresses gag, pol and env sequences is used to enable transfer of a secondary construct which carries an agronomically significant can and integration. Claimed isolated mucleic acid molecules comprise a nucleic acid sequence selected from a retroelement primer binding site, nucleic acid sequence selected from a retroelement primer binding site, envelope, gag, integrase, reverse transcriptase, protease or RNAse-H sequence (see AAZ5254-661). Also provided are plant retroviral particles that are used to transfer the nucleic acids into plant cells

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ABA82493;

Homo

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Human, mouse, Zmax1, HBM, high bone mass gene, lipid regulation, stroke, lipid-associated condition, arteriosclerosis, cardiovascular disease, ss, sosteoporosis, atherosclerosis, diabetic atherosclerosis, plaque build-up, neurovascular condition, wound healing, gene therapy, PCR primer, probe, bone development disorder, antiarteriosclerotic, cardiovascular,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a method for identifying a molecule involved in inhibits binding of a molecule to high bone mass (HBM) or its wild type gene, Zmaxi. Compounds identified by the method are useful for treating, diagnosing, preventing or screening for normal and abnormal lipidasses, stroke, and osteoporosis. The compounds may also be used in the treatment or prevention of diabetic atherosclerosis, cardiovascular treatment or prevention of diabetic atherosclerosis, neurovascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           conditions caused by plaque build-up, poor circulation due to plaque build-up and associated poor wound healing. The methods may be used in gene therapy, pharmaceutical development, and diagnostic assays for bone development disorders. Molecules identified by comparison of zmaxl and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HBM systems can be used as surrogate markers in pharmaceutical development, in diagnosis of human or animal bone disease, and in the treatment of bone disease. Sequences ABK22776-ABK23411 represent cDNA molecules encoding human Zmaxl and HBM, and PCR primers, probes, linkers and adapters of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Identifying molecules involved in lipid regulation, useful for diagnosing, treating or preventing e.g., arteriosclerosis, compidentifying a molecule that binds to high bone mass gene or its corresponding wild type gene.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seguence 18 BP; 4 A; 4 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Recker RR, Johnson ML;
                                                                                                                         Human Zmax1 cDNA reverse PCR primer #226.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 41; 409pp; English.
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                                                                                                                                                                                                                                                                                                                                  osteopathic; cerebroprotective
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es 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; high bone mass; HBM gene; Zmaxl gene; chromosome 11; 11q13.3; sequence tagged site; STS; osteoporosis; osteopathic; gene therapy; antisense therapy; vaccine; bone disorder; Paget's disease; adapter; sclerostosis; osteomalacia; fibrous dysplasia; PCK primer; linker; ss.
                                                                                                                         Gaps
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                                                         3.2%; Score 13.8; DB 1; Length 18; 88.2%; Pred. No. 2.3e+02; 1.ve 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Zmax1 gene region physical map preparation STS marker #452.
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Sequence 18 BP; 1 A; 4 C; 9 G; 4 T; 0 U; 0 Other;
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05-APR-2000; 2000US-00544398.
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Best Local Similarity 88.2.
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Query Match Best Loca Matches

RESULT 178

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ABK23290 ID ABK2 XX

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Neublastin DNA related PCR primer.
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Nootropic; neuroprotective; antiparkinsonian; anticonvulsant; analgesic; tranquiliser; antidiabetic; ophthalmological; neurodegenerative disorder; neublastin; ischemic neuronal damage; traumatic brain injury; diabetes; peripheral neuropathy; neuropatho; pain, Alzheimer's disease; glaucoma; Huntington's disease; Parkinson's disease; amyotrophic lateral sclerosis; memory impairment; renal disease; PCR; primer; 98.
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Unidentified.

WO200272826-A2.

19-SEP-2002.

12-MAR-2002; 2002WO-EP002691.

12-MAR-2001; 2001US-00804615.

(BIOJ ) BIOGEN INC. (NSGE-) NS GENE AS.

Rossomando Sah DWY, Johansen TE,

WPI; 2002-713515/77

New truncated neublastin polypeptides lacking one or more amino-terminal amino acids of a mature neublastin polypeptide useful for treating neurodegenerative disorders, e.g. peripheral neuropathy, neuropathic pain, brain injury.

Disclosure; Fig 8; 138pp; English.

The invention relates to a truncated neublastin polypeptide comprising an amino acid terminus that lacks one or more amino-terminal amino acids of a mature neublastin polypeptide. The polypeptides and nucleic acids are useful for treating neurodegenerative disorders such as ischemic neuronal damage, traumatic brain injury, peripheral neuropathy, neuropathic pain, Alzheimer's disease, Huntington's disease, Parkinson's disease, amyotrophic lateral sclerosis, memory impairment, disbetes, rend diseases, or glaucoma by moderating metabolism, growth, differentiation or survival of a nerve or neuronal cell. This polynucleotide sequence is a neublastin PCR primer of the invention

Sequence 18 BP; 3 A; 7 C; 6 G; 2 T; 0 U; 0 Other;

Gaps ; / Match 3.2%; Score 13.8; DB 1; Length 18; Local Similarity 88.2%; Pred. No. 2.3e+02; nes 15; Conservative 0; Mismatches 2; Indels

380 CCGCGACGACGGCGCCA 396 CTGCGACGACTGCGCCA 18

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RESULT 180 ACC45873

ACC45873 standard; DNA; 18 BP

ACC45873;

02-JUN-2003 (first entry)

Human HBM STS marker reverse primer #226.

Human; high bone mass; HBM; LRP5; LRP6; transgenic; bone mass modulation; gene therapy; bone dansity modulation; bone strangth; trabecular number; bone size; bone tissue connectivity; bone disease; osteoporosis; PCR; osteomalacia; rickets; Paget's disease; neoplasm of the bone; primer; ss.

sapiens Ното

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11-MAY-2001; 2001US-0290071P.
17-MAY-2001; 2001US-0291311P.
01-FEB-2002; 2002US-0353058P.
04-MAR-2002; 2002US-0361293P.
            13-MAY-2002; 2002WO-US014876
WO200292764-A2
      21-NOV-2002
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Babij P, Bex FJ, Yaworsky PJ, Bodine PV;

(GENO-) GENOME THERAPBUTICS CORP. (AMHP ) WYETH.

WPI; 2003-129278/12.

New transgenic animals (e.g. mice), useful as models for studying bone density modulation, developing drugs for treating or preventing bone diseases (e.g. osteoporosis), or diagnosing diseases characterized by reduced bone density.

Disclosure; Page 57; 603pp; English.

The invention relates to novel transgenic animals expressing the high bone mass (HBM) gene, expressing the corresponding wild type HBM gene, comprising an alteration of the gene encoding LRPS or LRP6, or expressing comprising an alteration of the gene encoding LRPS or LRP6, or expressing to make the study of bone density modulation. The transgenic animals are for the study of bone density modulation. The polymucleorides of the invention may have a use in gene therapy. The transgenic animals and muchical act of the study of bone density modulation, where the bone mass is modulated relative to non-transgenic animals of the same species in more than one parameter selected from bone density, bone size, or bone tissue connectivity. The transgenic animals, nucleic acids and methods are useful for identifying molecules involved in bone development, and for developing pharmaceutical compositions, which may be employed for treating or preventing bone diseases, e.g. osteoporosis, osteomalacia, rickets, Paget's disease, or neoplasms of the bone. The transgenic animals and nucleic acids are also useful in methods for disposing diseases in more development, or characterised by reduced bone density or mass. The present sequence is used in the exemplification of the invention

Sequence 18 BP; 4 A; 4 C; 7 G; 3 T; 0 U; 0 Other;

ö 3.2%; Score 13.8; DB 1; Length 18; 88.2%; Pred. No. 2.3e+02; tive 0; Mismatches 2; Indels Query Match Best Local Similarity 88.2 Matches 15, Conservative

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197 CTGCTCGGTGAAAGCAG 213 crecraeereacaecae 17

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ADB98571 standard; DNA; 18 RESULT 181 ADB98571 

ADB98571;

04-DEC-2003 (first entry)

Sequence tagged site #452 used to prepare Zmax1 (LRP5) gene region map.

Osteopathic; Gene therapy; High Bone Mass; HBM; LRP5; Zmax1; LRP6; bone mass modulation; osteoporosis; STS; sequence tagged site; ds.

Homo sapiens.

WO200292000-A2

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WPI; 1996-477128/47.
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                                                                                                                                                                                                                                                                                                The present invention relates to High Bone Mass (HBM), LRP5 (Zmax1) and LRP6 mutants, which results in a HBM-like phenotype when expressed in a cell. The HBM-like phenotype results in bone mass modulation and/or lipid level modulation. The invention is useful for diagnosting a HBM-like phenotype in a subject and for preparing a composition for modulating benoe mass and/or lipid levels in a subject suffering from e.g. osteoporosis. The present sequence is a Sequence Tagged Site (STS) marker, which was used to prepare a physical map of the Zmax1 (LRPS) gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lymphocyte specific interferon regulatory factor; LSIRF; IRF-3; probe; major histocompatibility complex; MHC; ISRE; interferon-stimulated response element; ds.
                                                                                                                                                                                                             New nucleic acid comprising a mutation in LRP5 or LRP6, useful for diagnosing a HBM-like phenotype in a subject and for preparing a composition for modulating bone mass and/or lipid levels in a subject
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                              Liu W;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                              Yaworsky PJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 18 BP; 4 A; 4 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
3.2%; Score 13.8; DB 1;
Best Local Similarity 88.2%; Pred. No. 2.3e+02;
Matches 15; Conservative 0; Mismatches 2;
                                                                                                                                                            Morales A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Grossman A, Richardson CD;
                                                                                                                                                              Graham JR,
                                                                                                                                                                                                                                                                          Example 2; Page 64; 629pp; English.
                                                                                                                                                                                                                                                   suffering from e.g. osteoporosis.
                                                                                                                       (GENO-) GENOME THERAPEUTICS CORP. (AMHP ) WYETH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       197 CIGCICGGIGAAGCAG 213
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11
                                                          11-MAY-2001, 2001US-0290071F.
17-MAY-2001, 2001US-0291311P.
04-PEB-2002; 2002US-0353058F.
04-MAR-2002; 2002US-0351293P.
                                    13-MAY-2002; 2002WO-US014877
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          96WO-CA000231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 95US-00422733.
96US-00611280.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAT41709 standard; cDNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 CTGCTAGGTGACAGCAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MHC ISRE binding sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CANADA INC
                                                                                                                                                            Allen K, Anisowicz A,
                                                                                                                                                                                      WPI; 2003-129214/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (AMGE-) AMGEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matsuyama T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9632477-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14-APR-1995;
03-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-JAN-1997
          21-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT41709;
                                                                                                                                                                                                                                                                                                                                                                                                        region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mus sp
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                                                                                      The murine major histocompability complex interferon-stimulated response element (MHC IRSE) binding sequence (AAT41709) was used as a probe to determine whether novel mouse lymphocyte-specific interferon regulatory factor (LSIRF) (see also AAR99426) is a DNA binding protein. LSIRF polypeptides were incubated with 32P- labelled double-stranded probe and, in some cases, with unlabelled competitor DNA fragments (see also AAT4170-16). Gel shift assays showed that the MHC ISRE sequence binds LSIRF protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                polymorphic genetic locus - used allele determination.
genes for murine lymphocyte specific interferon regulatory factor d for modulation of lymphocyte activation and proliferation.
                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 polymorphic; Human leukocyte antigen; HLA; DNA sequencing; PCR; polymerase chain reaction; allele; ss.
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                                                                                                                                                                                                                                                                                                                                       3.2%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 2.6e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3'-primer for HLA DR2 (15 and 16) allele amplification
                                                                                                                                                                                                                                                                                                 Sequence 19 BP; 7 A; 2 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Green RJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Identification of allele type of a known particularly for human leukocyte antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dunn JM, Leushner J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Page 17; 75pp; English.
                                                             Example 4; Page 40; 92pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                3 CAGAAGTGAAACTGAGG 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              95US-00577858.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (VISI-) VISIBLE GENETICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                  4 CAGGAGTGAAACTGCGG
                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 88.2
Matches 15; Conservative
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and evaluating four concurrent reactions, the sample is concurrently combined with at most three sequencing reaction mixtures containing different types of chain terminating nucleosides. The method can be used for the evaluation of polymorphic sites, and for determining the allelic type of a polymorphic gene. The methods are particularly useful for determining the HLA allele present in a sample
                                                                                                                                                                                                                                                                                                                                                                                                  Filamentous flower; FIL protein; agriculture; gardening; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This sequence represents a PCR primer for DNA encoding the filamentous flower (FIL) protein of the invention. The protein is useful in agriculture and gardening
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A gene participating in the flower formation of a plant useful in agriculture and gardening.
                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
3.2%; Score 13.8; DB 1; Length 19;
Best Local Similarity 88.2%; Pred. No. 2.6e+02;
Matches 15; Conservative 0; Mismatches 2; Indels
                                                                                                                           3.2%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 2.6e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Seguence 19 BP; 8 A; 5 C; 2 G; 4 T; 0 U; 0 Other;
                                                                                                 Sequence 19 BP; 2 A; 7 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                         PCR primer for FIL protein coding sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Page 7; 14pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      182 CAAGGCACATATCCACT 198
                                                                                                                                                                                      95
                                                                                                                                                                                                                                                                                      BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             98JP-00134095
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CAAGACACATATCAACT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAC73121 standard; DNA; 19
                                                                                                                                                                                      GCCGCGCAGTGGACATC
                                                                                                                                                                                                              18 GCGCGCGGTGGACACC
                                                                                                                                                                                                                                                                                    AAZ49122 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OKADA K.
MITSUI CHEM INC.
DAIICHI ENGEI KK.
TORAY IND INC.
                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                           15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-100767/09
                                                                                                                                           Sest Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                               Arabidopsis sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           JP11318462-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              L5-MAY-1998;
                                                                                                                                                                                                                                                                                                                                              06-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAC73121
                                                                                                                                                                                      79
                                                                                                                                                                                                                                                                                                                   AAZ49122;
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(DAII-) I
(TORA )
                                                                                                                               Query Match
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AAC73121/c
ID AAC731;
XX
AC AAC731;
                                                                                                                                                                                                                                                           RESULT 184
                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                        AAZ4912
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to an oligonucleotide array comprising oligonucleotide tags fixed to a solid substrate. The oligonucleotide array is useful for genotyping a nucleic acid sample at one or more loci array is useful for genotyping a nucleic acid sample a one or more loci anglify a polymorphic locus in a sample e.g. a single nucleotide polymorphism (SNP). The present sequence is one of the primers used in the method of the present invention to amplify a polymorphic sample amplified nucleic acid product is then used as a template in a SBE reaction with an extension primer. The SBE reaction products are used to
                                                     gonucleotide array, genotyping; single base extension reaction; SBE;
primer; polymorphic locus; single nucleotide polymorphism; ss.
                                                                                                                                                                                                                                                                                                                                                                                  Universal array of oligonucleotides tags attached to a solid substrate along with locus-specific tagged oligonucleotides useful in genotyping using single base extension reactions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pig; muscular steatosis-modulating factor; ss; metabolic; muscular; MS
food supplement; obesity; hyperlipidaemia; atherosclerosis;
wound healing; tumour; amyotrophic lateral sclerosis; ALS; PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                            Lockhart
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3.2%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 2.6e+02; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                            Lander ES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19 BP; 2 A; 7 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                              Kaplan P,
                                                                                                                                                                                                                                                                 (WHED ) WHITEHEAD INST BIOMEDICAL RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Porcine reverse PCR primer for TGFb.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 7; Page 49; 70pp; English.
                                                                                                                                                                                                                                                                                                              Huang X,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 282
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              form the oligonucleotide array
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        B
                                                                                                                                                                                         27-MAR-2000; 2000WO-US008069
                                                                                                                                                                                                                       99US-0126473P.
99US-0140359P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 266 GCACCTGGAGCAGGGCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18 GTACCTGGAGCAGAGCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAS62197 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                              Pan J, Hirschhorn JN,
Ryder T, Sklar P;
                                                                                                                                                                                                                                                                                   (AFFY-) AFFYMETRIX INC
                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-656171/63.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200179287-A2
                                                                                                                                 WO200058516-A2.
                                                                                                                                                                                                                       26-MAR-1999;
23-JUN-1999;
                                                                                                     Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-OCT-2001
                                                                                                                                                               05-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sus scrofa.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAS62197;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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Matches
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forward primer #13 used in multiplexing PCR/SBE assay.

(first entry)

02-FEB-2001

(CENE-) CENES PHARM INC.

WPI; 2002-097612/13.

Marchionni MA;

Page

Neuregulin-2 polypeptide and polynucleotide useful for treating multiple sclerosis, spinal muscular atrophy, nerve injury, Alzheimer's disease, by increasing mitogenesis, survival, growth or differentiation of a cell.

Example 1; Page 29; 79pp; English.

Wed Apr 21 12:58:21 2004

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The invention relates to prognosis or diagnosis of muscular steatosis by measuring the level of a muscular steatosis modulating factor (MSMF) in a human or animal and comparing this with the level in a healthy control. Any difference indicates presence of, or predisposition to, muscular steatosis. The method is particularly used for diagnosis or prognosis of muscular steatosis in mammals and birds, e.g. to select individuals as foundars in animal breeding. Also (ant)agonists of MSMF can be used to treat, or induce (for increasing the fat content of food) muscular steatosis, in humans and animals. The MSMF markers are also useful in the study of diseases and conditions such as obesity, hyperlipidaemla, atherosclerosis, wound healing, tumours and amyotrophic lateral sclerosis invention from its gene
                                                                                                                                                                                                     Prognosis and diagnosis of muscular steatosis, useful e.g. for selecting animals for breeding, by measuring levels of specific markers, also treating or inducing steatosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; ss; neuregulin-2; NRG-2alpha; NRG-2beta; mitogenesis; cell survival; cell growth; cell differentiation; erbB receptor; cardiomyopathy; ischemic damage; cardiac trama; heart failure; atherosclerosis; vascular lesion; vascular hypertension; 1531; degenerative congenital vascular disease; myasthenia gravis; neurodegenerative disorder; peripheral neuropathy; PCR primer; sensory nerve fiber neuropathy; motor fiber neuropathy; sensory nerve fiber neuropathy; multiple sclerosis; amyotrophic lateral sclerosis; spinal muscular atrophy; nerve injury; Alzheimer, disease; Parkinson's disease; cerebellar ataxia; spinal cord injury; tumour; neurofibromatosis; transgenic animal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3.2%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 2.6e+02; ative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19 BP; 4 A; 7 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                         (MIAC ) CANADA AGRIC & AGRI-FOOD CANADA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human Neuregulin-2 PCR primer 1531.
                                                                                                                                                                                                                                                                                     Example 1; Page 40; 190pp; English.
                                                                                                                              Gariepy C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAS18013 standard; DNA; 19 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               67 TGCACTACGAGGGCCGC 83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17 IGTACTACGIGGGCCGC 1
                                                     17-APR-2000; 2000US-0197936P.
                12-APR-2001; 2001WO-CA000509
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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Best Local Similarity 88.2 Matches 15; Conservative
                                                                                                                                                                       WPI; 2002-017600/02
                                                                                                                                Pomar C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAS18013;
                                                                                                                                Palin M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAS18013
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23-MAY-2001; 2001WO-US016896 23-MAY-2000; 2000US-0206495P.

WO200189568-A1. Homo sapiens

29-NOV-2001.

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The invention relates to a substantially pure neuregulin (NRG)-2

CD CD VRG-2beta (Cone 2DY) and the polymolectides encoding the Also

CC ON NRG-2beta (Cone 2DY) and the polymolectides encoding the Also

CC included are a vector expressing the protein, a host cell comprising the

CC included are a vector expressing the protein, a host cell comprising the

CC detains a knockout mutation in one or both NRG-2 allelses and an anti-NRG-2

CC having a knockout mutation in one or both NRG-2 allelses and an anti-NRG-2

CC diagnosing an increased likelihood of developing a NRG-2-related disease

CC diagnosing an increased likelihood of developing a NRG-2-related disease

CC diagnosing an increasing the

CC diagnosing an increasing the condition in a test subject. NRG-2 is useful for increasing the

CC mitogenels, survival, growth or differentiation of a cell (e.g. a

CC useful for treating diseases and disorders such as cardiomyopathy

CC useful for treating diseases and disorders such as cardiomyopathy

CC trauma or heart failure or which has a condition affecting smooth muscle

CC trauma or heart failure or which has a condition affecting smooth muscle

CC trauma or heart failure or which has a condition affecting smooth muscle

CC trauma or heart failure or which has a condition affecting smooth muscle

CC and degenerative congenital vascular lesion, vascular hypertension,

CC and degenerative disorder, peripheral neuropathy, a sensory nerve fiber

CC comprising inhibiting proliferation of a tumour cell, preferably a glial

CC comprising inhibiting proliferation of a tumour cell preferably a glial

CC comprising inhibiting proliferation of a tumour cell, correcting of neurofibromatosis by inhibiting glial cell

CC structure of cDNAs encoding NRG-2

CC structure of cDNAs encoding NRG-2

CC structure of connection of a tumour cell consideration cell consideration cell consideration cell considerat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human, single nucleotide polymorphism; nucleic acid typing; primer;
tissue typing; PCR; ACE; angiotensin coverting enzyme; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human angiotensin coverting enzyme SNP-fragment Bu6 PCR primer #1.
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3.2%; Score 13.8; DB 1; Length 19;
Best Local Similarity 88.2%; Pred. No. 2.6e+02;
Matches 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19 BP; 6 A; 7 C; 5 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /*tag= a
/note= "Biotinylated"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABN79916 standard; DNA; 19 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     45 GGCCACCACTCAGAGGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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modified_base
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Pourmand N;

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The invention relates to a novel method for obtaining typing information about several variable sites within target nucleic acid, or typing one or more nucleic acid molecules. The methods of the invention are useful for typing one or more nucleic acid molecules containing two or more variable sites, preferably nucleic acid molecules containing two or more variable variable sites are typed, where three or more primer extension reactions are performed. The method is also useful for diagnosis of pathological conditions characterized by the presence of specific nucleic acid molecule(s). The methods are particularly suited for identifying microbial species or this usbtypes, and in typing procedures e.g. typing of polymorphisms, tissue typing or in clinical applications. The sequence represents a PCR primer used in the invention to amplify a specific target region of genomic DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Adeno-associated virus; AAV; integration locus; CpG island;
SP1-like binding site; CAMP response element; CRE;
upstream binding factor 1; UBF-1; minisatellite; probe; gene therapy;
promoter; amplification; primer; polymerase chain reaction; PCR; ss.
                                                                                                                                                                                                              Typing nucleic acid for obtaining information about several variable sites involves simultaneously or sequentially performing two or more primer extension reactions, and determining the pattern of nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19 BP; 2 A; 8 C; 4 G; 5 T; 0 U; 0 Other;
                                                                           (PYRO-) PYROSEQUENCING AB.
(STRD ) UNIV LELAND STANFORD JUNIOR.
(GARD/) GARDNER R.
                                                                                                                                                                                                                                                                                                     Example 2; Page 47; 86pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              266 GCACCTGGAGCAGGGCG 282
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          10-SEP-2001; 2001WO-GB004042
                                             08-SEP-2000; 2000GB-00022069
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity 88.2
nes 15; Conservative
                                                                                                                                              Ronaghi M, Ekstroem B,
                                                                                                                                                                                   WPI; 2002-393849/42.
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                                                                                                                                                                                                                                                                    incorporation.
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18-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-SEP-1992;
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Matches
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                                                                                                                              In the cloning of AAVS1 from human lung fibroblast DNA, the primers given in AAQ63193-202 were used. A 4kb fragment contg. the AAV integration site was obtained (AAQ63192). (Updated on 25-MAR-2003 to correct PN field.)
                                               New nucleic acid corresponding to human adeno-associated virus integration site - useful e.g., as probe to confirm targetted integration of adeno-associated virus vectors in gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New HIV-type immune deficiency virus ECACC V 92092318 - and deriv. cDNA or antigens, useful for diagnosing retroviral infections and vaccines.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MVP-5180/91 DNA is obtained by PCR using the primers given in AAQ58925-
958. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-
2003 to correct PI field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human immunodeficiency virus; HIV; antigen; detection; diagnosis;
retrovirus; vaccine; lymphocyte; reverse transcriptase; amplification;
primer; polymerase chain reaction; PCR; 88.
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ilarity 88.2%; Pred. No. 2.9e+02;
Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                              3.2%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 2.9e+02; rative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hauser H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 5 A; 6 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                       Sequence 20 BP; 3 A; 6 C; 6 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eberle J, Brunn VA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 5; 73pp; German.
Linden RM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         92DE-04233646.
92DE-04235718.
92DE-04244541.
93DE-04318186.
                                                                                                        Claim 4; Page 4; 20pp; English.
                                                                                                                                                                                                                                                                       81 CGCGCAGTGGACATCAC 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               93EP-00116058
                                                                                                                                                                                                                                                                                            4
                                                                                                                                                                                                                                                                                          20 CGCTCAGAGGACATCAC
                                                                                                                                                                                                                                                                                                                                                                     AAO58941 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                         (revised)
(first entry)
                                                                                                                                                                                                                              Local Similarity 88.2 les 15; Conservative
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Best Local Similarity
Matches 15; Conserv
 Berns KI,
                        WPI; 1994-127741/16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  tat-1P primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22-OCT-1992;
30-DEC-1992;
01-JUN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Guertler LG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-OCT-1993;
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04-NOV-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                 Query Match
Best Local Si
Matches 15:
  Kotin RM,
                                                                                                                                                                                                                                                                                                                                                                                                 AAQ58941;
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Gaps ;

3.2%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 2.6e+02; tive 0; Mismatches 2; Indels

BP.

(revised)
(first entry)

19 GTACCTGGAGCAGAGCG 3

92US-00947127. 93EP-00114941.

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This invention describes the isolation of a novel HIV-type retrovirus called MVP-5180/91 (ECACC V 92092318). Antigens produced from this product can be used in an assay kit for detecting antibodies against viruses that cause immune deficiency, preferably where the assay is a Western blot, ELISA or fluorescence immunoassay. MVP-5180/91, cDNA and/or antigen can be used for detecting retroviruses that cause immune deficiency and to prepare vaccines. This sequence represents a PCR primer used in the method of the invention. (Updated on 20-MAR-2003 to correct PF field.) (Updated on 20-MAR-2003 to correct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Retinal calcium channel; RCC gene; alphalF-subunit; retinal disorder; myopia; nystagmus; strabismus; calcium-regulated development pathway; eye disorder; human; CACNAIF; CSNB; mutational analysis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                 New HIV-type retrovirus and corresponding cDNA, recombinant DNA and antigen - used for detecting retro-viruses that cause immune deficiency and to prepare vaccines.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Forward primer specific for human CACNAIF exon 16.
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88.2%; Pred. No. 2.9e+02;
tive 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                           Eberle J, Brunn AV, Knapp S,
                                                                                                                                                                                                                                                                                          (DADE-) DADE BEHRING MARBURG GMBH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 4; 39pp; German.
                   Synthetic.
Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       256
                                                                                                                                                                                               92DE-04235718.
92DE-04244541.
93DE-04318186.
93EP-00116058.
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                                                                                                                                             98EP-00114623
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nes 15; Conservative
                                                                                                                                                                                                                                                                                                                                                              WPI; 1999-072878/07.
                                                                                                                                                                                               22-OCT-1992;
30-DEC-1992;
01-JUN-1993;
05-OCT-1993;
                                                                                                                                                                                                                                                                                                                             Guertler LG,
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                                                                                                                                             05-OCT-1993;
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                                                                       EP890642-A2
                                                                                                        13-JAN-1999
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Best Local Si
Matches 15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Primers SS01 (given in AAQ76031) and SS02 (AAQ76032) were.used for the PCR amplification of a target region (AAQ76037) in the cytosine-DNA-methyltransferase of N. gonorrhoeae. Probe SS06-T5 (AAQ76033) is specific for a region in the amplified sequence, and is used to identify N. gonorrhoeae. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HIV-type retrovirus, MVP-5180/91, ECACC V 92092318; antigen, assay kit; detection; antibody, immune deficiency, vaccine; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Detection of Neisseria gonorrhoeae and/or Chlamydia trachomatis simultaneously by a simple, rapid and sensitive technique.
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3.2%; Score 13.8; DB 1; Length 20;
Best Local Similarity 88.2%; Pred. No. 2.9e+02;
Matches 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Seguence 20 BP; 5 A; 2 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                        Neisseria gonorrhoeae; probe; hybridization; cytosine-DNA-methyltransferase; CMT; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Fig 1; 29pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           367 TCACTITCCIGGACCGC 383
                   240 GGCTGCTTCCCGGGCTC 256
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                                                                                                                                             AAQ76033 standard; DNA; 20 BP.
                                      17 GGATGCTTCCAGGGCTC 1
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94US-00214861.
                                                                                                                                                                                                                                                                         N. gonorrhoeae probe SS06-T5
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16-JUL-1995
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19-MAY-1999
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AAX22342/
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Gaps .,

Length 20; 2; Indels

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Disclosure; Page 78; 115pp; English

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The invention provides a DNA molecule comprising a sequence of nucleotides encoding an alphalF-subunit of a mammalian retinal calcium channel (RCC), including a human alphalF-subunit, a murine alphalF-subunit and orthologs of the human and murine alphalF-subunits. The RCC gene may be used to develop products for diagnostic tests, for incomplete cSMB and risk assessment in affected families. The RCC gene can provide information as to the basic defect in this retinal conditions, which could lead to effective methods for treatment or cure of the disorder. As the associated features of myopia, nystagmus and strabismus frequently observed in patients with incomplete CSMB may be caused by calcium-requiated development pathways, identification of the RCC gene may help to elucidate the molecular details of eye development and which may lead to treatment for related eye disorders or diseases. Sequences AAZ46563-650 represent human CACNALF (alphalF-subunit of RCC gene) exon-specific CPCR primers, used for mutational analysis in humans
                                                                                                                                                                                                                                                                                                                              New isolated mammalian retinal calcium channel gene, used to develop products for the diagnosis and treatment of incomplete congenital stationary night blindness and related disorders.
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(UYTE-) UNIV TECHNOLOGIES INT INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure, Fig 6; 55pp; English
                                                                                                        Bech-Hansen T, Naylor MJ;
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Match 3.2%; Score 13.8; DB 1; Length 20; Local Similarity 88.2%; Pred. No. 2.9e+02; es 15; Conservative 0; Mismatches 2; Indels Query Match Matches

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AAZ44577 standard; DNA; 20 BP 07-APR-2000 (first entry) AAZ44577; 

Newcastle disease virus LaSota primer p1898-.

Avian-paramyxovirus; infection; lentogenic; F protein; vaccine; respiratory disease; gastrointestinal disease; poultry pathogen; local immunity; primer; ss.

Newcastle disease virus

23-DEC-1999.

WO9966045-A1.

99WO-NL000377. ,7-JUN-1999; 19-JUN-1998; DIEN-) STICHTING DIENST LANDBOUWKUNDIG ONDERZOE.

98EP-00202054.

Gielkens ALJ; De Leeuw OS, Koch G, Peeters BPH,

WPI; 2000-106102/09.

New avian paramyxovirus cDNA, useful for production of vaccine against Newcastle disease virus.

This invention describes a novel avian-paramyxovirus cDNA (I) which comprises a nucleic acid sequence corresponding to the 5' terminal end of the genome of avian-paramyxovirus allowing the generation of an infectious copy of avian-paramyxovirus. The cell line is useful for the production of infectious lentogenic NDV (Newcastle Disease virus) without the addition of exagence proteolytic activity. Also it is possible to generate a stable transfected cell line that expresses the wild-type F protein in the virus envelope therefore providing infectious particles, useful in the form of a vaccine, especially adding respiratory and/or gastrointestinal diseases. NDV can be easily cultured to very high titers in embryonated eggs. Mass culture of embryonated eggs is relatively cheap. NDV vaccines are relatively stable and can be simply adminiscered by mass application methode e.g. darinking water or by spraying or by aerosol formation. The natural route of infection is by the respiratory cand/or gastrointestinal tract which are also the mady or routes of infection of many other poultry pathogens. NDV can induce local immunity despite the presence of circulating maternal antibody. AAZ44512-Z44609 and AAZ44618-Z44560 represent primers used in the isolation of the NDV strain Lascha cannones. strain LaSota genome 

Sequence 20 BP; 4 A; 4 C; 6 G; 6 T; 0 U; 0 Other;

Gaps ô / Match 3.2%; Score 13.8; DB 1; Length 20; Local Similarity 88.2%; Pred. No. 2.9e+02; les 15; Conservative 0; Mismatches 2; Indels Query Match Best Local

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5717/c AAD06717 standard; DNA; 20 BP. 10-AUG-2001 (first entry) AAD06717; 

RESULT 195

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Gaps

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Phenylalanyl-tRNA synthetase; PheRS; amino acid separation; ATP quantitation; protein inhibitor; antimicrobial; antibiotic effect; PCR primer; ss. C-terminal phenylalanyl-tRNA synthetase DNA amplifying primer, EfP-5.

Enterococcus faecalis.

US6221640-B1.

24-APR-2001.

97US-00855910. 14-MAY-1997;

14-MAY-1997; 97US-00855910.

(CUBI-) CUBIST PHARM INC.

Avruch AS, Shen X, Sassanfar M, Gallant PL, Tao J, Nair S;

Yu RV;

WPI; 2001-327244/34.

New Enterococcus faecalis aminoacyl-tRNA synthetase proteins and nucleic acids useful for separating amino acids which they specifically recognize, in quantifying amino acids and ATP, or for detecting protein inhibitors.

Example 3; Col 41; 88pp; English.

The present invention relates to Enterococcus faecalis aminoacyl-tRNA synthetases are useful in the biochemical

Page

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separation of the amino acid which they specifically recognise and in identifying inhibitors ocid and APP, and for detecting and the amino acid and APP. And for detecting and these enzymes can be because the potential inhibitors of these enzymes can be screened for antimicrobial or antibiotic effects, without requiring the culture of pathogenic strains of Enterococcus. The purification and study of the enzymes can be made and used in the purification of protectine or to Diypeptides, and the aninoacyl-tRNA synthetase genes may be used as probes to identify DNA fragments encoding the corresponding aminoacyl-tRNA synthetase gene from other species of enterococci by specific hybridisation. The present sequence is a PCR primer which is used for amplifying the C-terminal Enterococcus facecalis phenylalanyl-tRNA synthetase (PherS) DNA
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Sequence 20 BP; 5 A; 6 C; 7 G; 2 T; 0 U; 0 Other;

ö 0; Gaps 3.2%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 2.9e+02; ative 0; Mismatches 2; Indels Query Match Best Local Similarity 88.2 Matches 15; Conservative

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AAS09653/c 

AAS09653 standard; DNA; 20 BP.

AAS09653;

(first entry) 26-SEP-2001 Immunoreactive CpG sequence-containing oligonucleotide #103.

CpG sequence; immune response; non-B cell activation; interferon gamma; IFN-gamma; humoral; antibody production; interleukin-6 production; herapeutic; allergy; asthma; cancer; autoimmune disorder; infection; bio-warfare, vaccine; antisense therapy; eczema; allergic rhinitis; coryza; hay fever; urticaria; hives; food allergy; atopic condition; hepatitis; human immunodeficiency virus; HIV; malaria; Francisella; lupus erythematosus; rheumatoid arthritis; multiple sclerosis; schistcoschiasis; tuberculosis; acquired immunodeficiency syndrome; AIDS; leishmania; Ebola; Anthrax; Listeria; ss.

Synthetic

WO200151500-A1

19-JUL-2001

12-JAN-2001; 2001WO-US001122

(USSH ) US DEPT HEALTH & HUMAN SERVICES.

14-JAN-2000; 2000US-0176115P.

Verthelyi D;

Ishii K, Klinman D,

WPI; 2001-442129/47.

Oligodeoxynucleotides for inducing an immune response to treat and prevent an allergic reaction, cancer, an autoimmune disorder and symptoms resulting from exposure to bio-warfare agents, comprise multiple CpG AASO9551-AASO9662 represent oligodeoxynuclectides (ODN) of at least 10 nuclectides comprising multiple CpG sequences, where one of the CpG sequences is different from another of the multiple CpG sequences. The ODN are useful for inducing an immune response, preferably a cell-mediated immune response, preferably a cell-mediated immune response, involving non-B cell activation, interferon Claim 5; Page 44; 48pp; English. sednences

cell activation, antibody and interleukin-6 production in a host, for treating, preventing or ameliorating an allergic reaction, e.g. asthma, cancer, e.g. solid tunnour cancer, a disease associated with the immune cancer, e.g. solid tunnour cancer, a disease associated with the immune system e.g. autoimmune disorder or an immune system deficiency, infection or a symptom resulting from exposure to bio-warfare agent in a human. The condition of immune response improves the efficacy of a vaccine and is used in antisense therapy. The OND are useful for treating preventing cused in antisense therapy. The OND are useful for treating preventing or ameliorating allergic reactions, including eczema, allergic rhinitis or coryza, hay fever, bronchial asthma, untitarial (hives), food allergies and other atopic conditions, for improving the efficacy of vaccines against hepatitis A, B and C, human immunodeficiency virus (HIV) and calaria, for treating immune system deficiencies, e.g. lupus cervitations and autoimmune diseases such as rheumatoid arthritis and multiple sclerosis, infections including Francisella, schistosomiasis, tuberculosis, acquired immunodeficiency syndrome (AIDS), Leishmania and symptoms resulting from exposure of bio-warfare agent, including Ebola, conditions and Listeria response involving B

Sequence 20 BP; 2 A; 4 C; 12 G; 2 T; 0 U; 0 Other;

Gaps ö Query Match
3.2%; Score 13.8; DB 1; Length 20;
Best Local Similarity 88.2%; Pred. No. 2.96+02;
Matches 15; Conservative 0; Mismatches 2; Indels

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RESULT 197 AAS21720/c

AAS21720 standard; DNA; 20

Bb

AAS21720; 

(first entry) 21-NOV-2001 Mouse Survivin antisense oligonucleotide #23.

Survivin; human; mouse; cytostatic; antisense oligonucleotide; hyperproliferative condition; cancer; apoptosis; cytokinesis;

Mus musculus. Synthetic.

WO200157059-A1.

09-AUG-2001.

30-JAN-2001; 2001WO-US002939.

02-FEB-2000; 2000US-00496694.

(ISIS-) ISIS PHARM INC

Swayze EE, CF, Ackermann EJ, Bennett

Cowsert LM;

WPI; 2001-488863/53.

Novel antisense compounds for modulating the expression of Survivin and treatment of cancer.

Example 18; Page 60; 120pp; English.

The invention relates to antisense oligonucleotides targeted to a nucleic acid molecule encoding human Survivin, where the antisense oligonucleotide inhibits the expression of human Survivin. These antisense oligonucleotides are used in the treatment of an animal suffering from a disease or condition associated with Survivin, e.g. a hyperproliferative condition such as cancer, and comprises administering a therapeutically or prophylactically effective amount of the antisense oligonucleotide so that expression of Survivin is inhibited. The

Candida albicans GRACE strain PCR primer SEQ ID NO 4516.

(first entry)

30-JAN-2003

ABZ30365;

ABZ30365 standard; DNA; 20 BP.

ABZ30365

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oligonucleotides can also be used to treat a human suffering from a disease or condition characterised by a reduction in apoptosis comprising administering the antisense oligonucleotide to a human. In addition, the antisense oligonucleotide and a cytotoxic chemotherapeutic agent e.g. taxol or cisplatin, can be used to modulate apoptosis, cytokinesis or the cell cycle, or inhibit the proliferation in a cancer cell by contacting the cell with the antisense oligonucleotide. AAS21521-AAS21768 represent Survivin nucleic acids, and antisense oligonucleotides targeted to Survivin, used in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention provides a microarray of oligonuclectides comprising probes for the human HLA Class I genes attached to a solid support. These can be used in HLA typing. Oligonuclectide arrays are also useful in large scale gene discovery, monitoring gene expression, polymorphism detection and gene mapping
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oligonucleotide arrays useful for human leukocyte antigen (HLA) tissue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             typing, comprises HLA class I oligonuclectide probes representing all known polymorphisms in HLA class I locus, on a solid support.
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                                                                                                                                                                                                                       3.2%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 2.9e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human HLA Class I oligonuclectide probe SEQ ID NO: 38.
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(UNIW ) UNIV WASHINGTON.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure, Page 54; 83pp; English.
                                                                                                                                                                                                                                                                                                358 GCGACTTCCTCACTTTC 374
                                                                                                                                                                                                                                                                                                                                                                                                                             AAF54593 standard; DNA; 20 BP.
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Best Local Similarity
Matches 15; Conserva
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The invention relates to constructing (M1) a strain of diploid fungal cells in which both alleles of a gene are modified, comprising modifying cells in which both alleles of a gene are modified, comprising modifying cone allele by insertion or teplacement fragment with a heterologous promoter, so that expression of the second allele is regulated by the promoter, so that expression of the second allele is regulated by the promoter. (M1) is useful for constructing a strain of diploid fungal cells in which both alleles of a gene are modified. The diploid fungal cells in which both alleles of a gene are useful for identifying a gene that is essential to the survival or growth of a fungus, a gene that is essential to the survival or growth of a fungus, a gene that contributes to the resistance of a diploid fungus to an antifungal agent that inhibits the growth of a mammalian disease. (M1) is useful for identifying a compound which modulates the compound catabolism, biosynthetic, transporter, transcriptional, compound extabolism, biosynthetic, transporter, transcriptional, transduction, DNA replication and cell division compound stabolism, biosynthetic, transporter, transcriptional, transduction, both relativing a compound having the ability to inhibit growth of relativing a compound having the ability to inhibit growth or proliferation of C. albicans cells and for treating infection by C. albicans. The present sequence is that of the primer used in the method of the invention. Note: The sequence data for this patent is not represented in the printed specification but is based on sequence information supplied to Derwent by the European Patent Office
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Constructing strains for identifying gene products as effective targets for therapeutic intervention, by inactivating in the strain one allele of a gene and placing other allele of the gene under conditional expression.
                                                                                                                                                                        Fungus; yeast; tetracyclin; promoter; GRACE strain; biosynthesis;
signal transduction; DNA replication; cell division; growth;
proliferation; Candida albicans; fungicide; antifungal; PCR; primer; ss.
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18.2%; Pred, No. 2.9e+02;
ve 0; Mismatches 2; Indels
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22-AUG-2001; 2001US-0314050P.
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les 15; Conservative
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                                                                                                                                                                                                                                                                 Candida albicans.
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Gaps

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2; Indels

0; Mismatches

298 AGGACCTGAGCCCCGGG 314

15; Conservative

Matches

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Similarity

Query Match Local AGGACCTGAGCTCCTGG 18

3.2%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 2.9e+02;

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The invention relates to constructing (M1) a strain of diploid fungal cells in which both alleles of a gene are modified, comprising modifying cells in which both alleles of a gene are modified, comprising modifying one allele by insertion or treplacement fragment with a heterologous promoter, so that expression of the second allele is regulated by the promoter. (M1) is useful for constructing a strain of diploid fungal cells in which both alleles of a gene are useful for identifying a gene that cells in which both alleles of a gene are useful for identifying a gene that is essential to the survival or growth of a fungus, a gene that is essential to the survival or growth of a fungus, a gene that contributes to the virulence and/or pathogenicity of a fungus to an antifungal agent that inhibits the growth of a diploid fungus to an antifungal agent that inhibits the growth of a mammalian disease. (M1) is useful for identifying a compound which modulates the activity of a gene product, preferably enzymatic activity, carbon compound catabolism, biosynthetic, transporter, transcriptional, translational, signal transduction, DNX replication and cell division activity. The method is useful for identifying a compound having the ability to inhibit growth or proliferation of C. albicans cells and for treatment is not represented in the present sequence at for present cell in the Buropean Patent Office on sequence information supplied to Derwent by the Ruropean Patent Office
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Constructing strains for identifying gene products as effective targets for therapeutic intervention, by inactivating in the strain one allele of a gene and placing other allele of the gene under conditional expression.
                                                                                                                                                                                                                                                                                                                                                                                                                              Fungus; yeast; tetracyclin; promoter; GRACE strain; biosynthesis;
signal transduction; DNA replication; cell division; growth;
proliferation; Candida albicans; fungicide; antifungal; PCR; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                 Candida albicans GRACE strain PCR primer SEQ ID NO 5310.
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                                                                             ABZ31091 standard; DNA; 20 BP
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22-AUG-2001; 2001US-0314050P.
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                                                                                                                                                                                                                                                          30-JAN-2003 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Candida albicans.
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                                                                                                                                                                        ABZ31091;
RESULT 200
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ö 0; Gabs Query Match 3.2%; Score 13.8; DB 1; Length 20; Best Local Similarity 88.2%; Pred. No. 2.9e+02; Matches 15; Conservative 0; Mismatches 2; Indels

1 GCCAAATCGGAAGACTG 17 셤 AAD45182 standard; DNA; 20 BP

AAD45182,

Human RIP2 antisense oligonucleotide ISIS #104252.

27-DEC-2002 (first entry)

AAD45182;

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New antisense oligonucleotide that targets regions of a nucleic acid encoding human receptor interacting protein (RIP)2, for treating diseases associated with RIP2 expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to antisense compounds targetted to a nucleic acid encoding human receptor interacting protein (RIP)2 to inhibit its expression. Antisense compounds are used for treating diseases associated with RIP2 expression. They are also useful in antisense gene therapy. The present sequence is an oligonucleotide targetted to human RIP2 DNA
Human, receptor interacting protein, RIP2, antisense, gene therapy, phosphorothicate, ss.
                                                                                                                                                                                                                                                                                                                  note = "2-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    note= "2-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                        note= "Phosphorothioate backbone"
                                                                                                                                     Location/Qualifiers
                                                                                                                                                             *tag= c
mod_base= OTHER
                                                                                                                                                                                                                                                                                             'mod_base= OTHER
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/mod_base= m5c
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/mod_base= m5c
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/mod_base= m5c
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                                                                                                                                                                                                                                                                                                                                                             *tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-673017/72.
                                                                                                                                       Key
modified_base
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                                                                  Homo sapiens.
Synthetic.
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228 GCCAAATCGGGAGGCTG 244

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Gaps

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The invention relates to a novel isolated nucleic acid molecule comprising a sequence that encodes a thioesterase or thioesterase domain, derived from a bacterial daptomycin bloopyntheric gene duster. The proteins of the invention have antibacterial, fungicide, virucide, antiparasitic, immunomodulator, antibacterial, fungicide, virucide, compositions and have a use in gene therapy. The compositions and methods of the present invention are useful for generating novel linear and cyclic peptides and improving yield of a product in a cell expressing an daptomycin non-ribosomal peptide synthetase (NRFS) to be used as new compounds or in producing new compounds, such as antibiotics, antitumour agents, antifungals, antivirals, antiparasilics, antimitotics, antitumour agents, immunomodulatory agents, anti-cholesterolemic agents, siderophores, approachemicals and synostatics. The sequence represents a PCR primer used in the invention to amplify the S. roseosporus daptomycin blosynthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Isolated nucleic acid molecule from a bacterial daptomycin biosynthetic gene cluster encoding a thioesterase or thioesterase domain, useful for generating novel linear and cyclic peptides, and products in a cell.
                                                                                                                                                                                                                                                                                                                                                                                                         Daptomycin biosynthetic gene cluster; thioesterase; antibacterial; fungicide; virucide; antiparasitic; immunomodulator; antilipemic; cytostatic; gene therapy; antimitotic; immunomodulatory; siderophore; anti-cholesterolemic; agrochemical; linker; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                       S. roseosporus daptomycin biosynthetic gene cluster PCR primer P76
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                                             / Match 3.2%; Score 13.8; DB 1; Length 20; Local Similarity 88.2%; Pred. No. 2.9e+02; les 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Seguence 20 BP; 4 A; 8 C; 5 G; 3 T; 0 U; 0 Other;
             Sequence 20 BP; 2 A; 7 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Silva CJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 2; Page 91; 227pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Baltz RH,
                                                                                                                             CCTGAGCCCCGGGGACC 318
                                                                                                                                                                                                                                                             ABQ78909 standard; DNA; 20 BP.
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28-FEB-2001; 2001US-0272207P.
06-AUG-2001; 2001US-0310385P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Streptomyces roseosporus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-599794/64
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BALTZ R H.
SILVA C J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200259322-A2
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(BALT/)
(SILV/)
                                                     Query Match
                                                                                                                                                                                                                           RESULT 202
                                                                                         Matches
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DB 1; Length 20;

3.2%; Score 13.8;

Query Match

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The present invention describes a method (M1) for designing capture oligonuclectide probes (II) will hybridise with little mismatch, where cligonuclectide probes (II) will hybridise with little mismatch, where coligonuclectide probes (II) will hybridise with little mismatch, where coligonuclectide probes (II) will hybridise with little mismatch, where coligonuclections deserved within a narrow range. The method is useful infectious agents e.g. Cryptococcus neoformans, Candida albicans and infectious agents e.g. Cryptococcus neoformans, Candida albicans and Aspergillus fumigautus, viruses e.g. T-cell lymphocytotrophis cirus, Bestein-Barr virus and polio virus, and parabitic infectious agents confined is also useful for detecting genetic diseases such as 21 hydroxylase deficiency, Turner Syndrome and obesity defects.

Confecting cancer involving oncogenes, tumour suppressor genes, or genes involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCAl gene, psi gene, human papilinaavirus types 16 and 18 and liver cancers. The cancer is also used for environmental monitoring, forensics and the food and feed industry, detecting comprises scanning (using e.g. a scanning electron microscope and infrared microscope) the support at the control problem of the oligonucleotide probe presence or absence of the target uncleotide sequences. All SO74 to a presence or absence of the target uncleotide sequences. All so in a contraction of the oligonucleotide problem.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 e of the target nucleotide sequences. ABI82074 to oligonucleotide sequences used in the exemplification
                                                                                                                                                                                                                                                                                                                                                  Human, K-ras, PCR primer; probe; capture probe; mutation detection; ligase detection reaction; LDR; p53; BRCA1; BRCA2; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndrome; obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forenaic; environmental monitoring; food industry; feed industry; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Designing capture oligonucleotide probes for use on a support to which complementary oligonucleotides hybridize with little mismatch.
                   Gape
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                   Indels
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88.2%; Pred. No. 2.9e+02;
ive 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                  Capture oligonucleptide Zip ID#1931 oligo #9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 13.8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 5; Fig 29; 300pp; English.
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                                                           373 TCCTGGACCGCGACGAC 389
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                                                                                                                                                                                                    BP.
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                                                                                                 TACTGGACCTCGACGAC
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                     15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-034366/04.
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Best Local Similarity
Matches 15; Conserv
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                                                                                                                                                                                                                                                                                16-FEB-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                        ABI94844;
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ABI94844/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation coodon, coding regalon, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 mucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dystunction and a second active agent comprising an entilnflammatory steroid and ubjquinone. A composition of the invention has antiinflammatory, antiallargic, antiathmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an application of or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubjquinone or lung surfactuant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.
                                                                                                                                                                                                                                                                    Human, antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pabalan J, Aguilar D;
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               Indels
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Pred. No. 2.9e+02;
0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                               lung inflammation; respiratory disease; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Katz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 15; SEQ ID NO 447; 872pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Li Y, Sandrasagra A, Ka
Tang L, Shahabuddin S;
 Best Local Similarity 88.2st_i, Promatches 15, Conservative 0;
                                                                                                                                                                                                                                           Human oligonucleotide sequence
                                            345 CGGCTGCTCTACAGCGA 361
                                                                                                                                                 ВP
                                                                       18 cggcrgcgaracagcga 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                    ABZ85205/c
ID ABZ85205 standard; DNA; 20
                                                                                                                                                                                                           (first entry)
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                                                                                                                                                                                                            17-0CT-2003
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Miller S,
                                                                                                                                                                                 ABZ85205;
                                                                                                                      RESULT 204
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The present invention describes a method and a kit for determining the expression of mRNA or cDNA of a protein participating in the maintenance of skin structure. The method is quantitative, simple and accurate in the determination of extracellular matrix components of laminin 5 chain genes LAMA3, LAMB3 and LAMC2, matrix metalloproteinases MMP-1, MMP-2, MMP-3 and MMP-9, VII collagen, type I collagen alpha 1 chain, type I collagen alpha 1 chain, type IV collagen alpha 1 chain, type IV collagen alpha 2 chain, TIMP-1, TMMP-3 and TIMP-3. ACF57290 represent PCR primers and probes used in the method of the
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; mouse; skin structure; skin; laminin 5 chain gene; LAMA3; LAMB3; LAMC2; extracellular matrix component; matrix metalloproteinase; MMP-1; MMP-2; MMP-3; MMP-3; MMP-9; TIMP-3; TIMP-3; collagen; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A method and a kit for determination of expression of mRNA or cDNA of protein participating in the maintenance of skin structure.
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88.2%; Pred. No. 2.9e+02;
tive 0; Mismatches 2; Indels
                            Indels
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  88.2%; Pred. No. 2.9e+02;
ive 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                       Human TIMP-2 reverse PCR primer SEQ ID NO:83.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 4; 34pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           144 GCGGTGGAGGCCGGCTT 160
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                                                                        245 CITCCCGGGCTCGGCCA 261
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                                                                                                                         CTACCAGGGCTCGGCCA
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Best Local Similarity 88.2
Matches 15; Conservative
                            Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-407328/39.
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JP2002330792-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
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                                                                                                                                                                                                                                                                                             ACF57283;
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                                                                                                                                                                                                 RESULT 205
ACF57283
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Length 20;

DB 1;

Score 13.8;

Query Match

Antinsense oligonucleotide targeting mouse C3 component, ISIS140049

04-DEC-2003 (first entry)

wouse; ss; antisense; complement component C3; inflammation; septic shock; multiple organ failure; hyperacute organ failure; autoimmune disorder; CNS inflammation; multiple sclerosis; atherosclerosis; tumour.

mod_base= OTHER
note= "Phosphorothioate backbone and all cytosines are
nethyl cytosines"

Location/Qualifiers

Key modified_base

Mus musculus

Д

/mod_base= OTHER /note= "2'-methoxyethyl nucleotides" 16. .20

*tag= a

modified_base

/mod_base= OTHER /note= "2'-methoxyethyl nucleotides"

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*tag=

modified base

23-OCT-2001; 2001US-00001076. 23-OCT-2001; 2001US-00001076.

US2003096775-A1

22-MAY-2003

(ISIS-) ISIS PHARM INC

Watt AT;

Graham MJ,

WPI; 2003-606441/57.

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New compound, having a sequence targeted to a nucleic acid encoding mucin 1, transmembrane, useful for preparing a composition for treating hyperproliferative or inflammatory disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to antisense oligonucleotides targetted to a nucleic acid encoding mucin 1 transmembrane (also known as WCT, epistalin, epitectin, polymorphic epithelial mucin; PEM, peanut-reactive urinary mucin; PUM, epithelial membrane antigen, BEM, PAS-0, NCRC11, H23 antigen, H23-EFA transmembrane antigen, PFB antigen and CD227) to inhibit/modulate the expression of mucin 1 transmembrane. Antisense compounds of the invention are useful for preparing compositions for transmet hyperproliferative or inflammatory disorders. The invention is also used in gene therapy. The present sequence is human mucin 1 transmembrane antisense oligonucleotide
          Human, mucin 1 transmembrane; hyperproliferative disorder; cytostatic; inflammatory disorder; gene therapy; H23-ETA transmembrane antigen; antisense; episialin; epitectin; polymorphic epithelial mucin; CD227; peanut-reactive urinary mucin; PUM; epithelial membrane antigen; BEM; PEM; NCRC11; H23 antigen; DF3 antigen; phosphorothioate backbone; MUC1;
                                                                                                                                                                                                              /mod_base= OTHER
/note= "Phosphorothicate backbone; All cytidines are 5-
methyl cytidines"
                                                                                                                                                                                                                                                                                                                                                                      /mod_base= OTHER
/note= "2'-methoxyethoxy (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                          mod base= OTHER
'note= "2'-methoxyethoxy (2'-MOE) nucleotides"
16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 4 A; 3 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 3; Page 82; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20-DEC-2001; 2001US-00029517.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-DEC-2002; 2002WO-US039873.
                                                                                                                                                                                                                                                                            ...5
/*tag= b
                                                                                                                                                                                                                                                                                                                                                        *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Myers SJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-559135/52.
                                                                                                                                                                                                                                                                                                                                                                                                                         WO2003054154-A2
                                                                                                                                                                           Key
modified_base
                                                                                                                                                                                                                                                                          modified base
                                                                                                                                                                                                                                                                                                                                       modified_base
                                                                                                                           sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                        03-JUL-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Dobie KW,
                                                                                                                            Homo sapie:
Synthetic.
                                                                                           PAS-0; SB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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Gaps
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                                                                        3.2%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 2.9e+02; ive 0; Mismatches 2; Indels
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247

231 AAATCGGGAGGCTGCTT

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Local Similarity

Best Loc Matches

20

ATATCGAGAGGCTGCTT

B5.

ADB89961 standard; DNA; 20

RESULT 207

ADB89961

ADB89961;

SXB

New antisense oligonucleotides targeted to a nucleic acid molecule encoding complement component C3, useful for treating a disease or condition associated with complement component C3, e.g. autoimmune

Claim 3; Page 27; 72pp; English.

disorder or infection.

The invention relates to a compound 8-50 nucleobases in length targeted to a nucleic acid molecule encoding complement component C3. The compound complement component C3 and inhibite the expression of complement component C3 or specifically hybridises with at least an 8-nucleobase component C3, or specifically hybridises with at least an 8-nucleobase component C3. Also included are a composition comprising complement component C3. Also included are a composition comprising the component C3 in calls or tissues (comprising the expression of complement component C3 in calls or tissues (comprising contacting the cclls or tissues with the compound cited above) and treating an animal compounds are useful for inhibiting the expression of complement component C3 comprising deministering to the animal the compound cited above so that expression of complement component C3 is inhibited. The antisense compounds are useful for inhibiting the expression of complement c3 in cells or tissues, or for treating an animal having a disease or condition associated with complement component C3 such as an expression of complement c3 in cells or tissues, or for treating an animal having a disease or condition associated with complement component C3 such as an expression of capan failure, component C3 in cells or tissues, or for treating an animal having a disease or condition associated with complement component C3 such as an expression of capan failure and C4 of various members of a biological pathway, or for preventing or delaying the expression of infection, inflammation or tumour formation. The componence is an expression of infection or tumour formation. The present sequence is an expression or income capan pathway, or for preventing or delaying an expression or income capan path and delaying and expression or tumour formation. The present sequence is an expression or income capan path and Sequence 20 BP; 4 A; 9 C; 3 G; 4 T; 0 U; 0 Other; antisense oligonucleotide targeting mouse C3.

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The present invention describes a method for inducing the production of vascular endothelial growth factor (VEGF) by a cell comprising contacting the cell with a CpG oligonucleotide and therefore inducing the production of VEGF by the cell. Also described: (1) inducing neovascularisation in the cell into an area of the tissue, comprising introducing a CpG oligonucleotide into an area of the inducing neovascularisation in the whole wessels is desired, and so inducing neovascularisation in the subject where angiogenesis is desired, comprising introducing a CpG oligonucleotide to the area, and so promoting angiogenesis in the subject where angiogenesis is desired, comprising introducing a CpG oligonucleotide to the area, and so promoting angiogenesis in the subject; and (3) screening for an agent that inhibits neovascularisation, comprising administering the agent to command, where inhibition of angiogenesis in the animal indicates that the mammal, where inhibition of angiogenesis in the animal indicates that the captorides can be used in gene therapy. The method and the CpG oligonucleotides can be used in inducing angiogenesis or coligonucleotides can be used in inducing angiogenesis or exhibit male pattern baldness, or subjects who have a wound or who have constituted and the inhibit neovascularisation. The present sequence for agents that inhibit neovascularisation. The present sequence
                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Inducing the production of vascular endothelial growth factor by a cell, useful for inducing angiogenesis, comprises contacting the cell with a CpG oligodeoxynucleotide.
                                                                                                                                                                                                                                                                                                                                                        vascular endothelial growth factor; VEGF; CpG oligonucleotide; neovascularisation; angiogenesis; vulnerary; vasotropic; antiarteriosclerotic; gene therapy; skin graft; male pattern baldness; atherosclerosis; ischaemia; ss.
                                       Gaps
                                       ö
Score 13.8; DB 1; Length 20;
Pred. No. 2.9e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (UYTE-) UNIV TENNESSEE RES CORP.
(USSH ) US DEPT HEALTH & HUMAN SERVICES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 7; SEQ ID NO 45; 37pp; English.
                                                                                                                                                                                                                                                                                                                          CpG D oligonucleotide SEQ ID NO:45.
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                                                                                                                                                                                                           ADD01081 standard; DNA; 20 BP
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   3.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-DEC-2002; 2002WO-US040955.
                                                                             38 CGAAGATGGCCACCACT
                                                                                                               ccaactroccact
                                                                                                                                                                                                                                                                                     (first entry)
                     Similarity 88.2
5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Klinman DM, Zheng M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-559138/52.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO2003054161-A2.
                                                                                                                                                                                                                                                                                     01-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-JUL-2003
                                         15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                   ADD01081;
   Query Match
Best Local 6
                                         Matches
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Sequence 20 BP; 2 A; 4 C; 12 G; 2 T; 0 U; 0 Other;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Detection of herpes B virus by PCR amplification of sample DNA - to detect a specific herpes simian monkey B virus DNA segment.
                                                                                                                                                                                                                                              Primer; polymerase chain reaction; PCR; diagnosis; herpes B virus; primate alpha-herpes virus gB glycoprotein; ss.
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                          ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 21;
Length 20;
                                                                                                                                                                                                                        B3' for primate alpha-herpes gB glycoprotein.
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                          Indels
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Score 13.8; DB 1; ... Pred. No. 2.9e+02; ... Mismatches 2;
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88.2%; Pred. No. 3.2e+02;
iive 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                Black
                                                                                                                                                                                                                                                                                                                                                                                                                                                Eberle R,
                                                                                                                                                                                                                                                                                                                                                                                                                       (SWBI-) SOUTHWEST FOUND BIOMEDICAL RES.
                          0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Scinicariello F,
                                                      254 CTCGGCCACGGTGCACC 270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              38
                                                                                                                                              BP
                                                                                                                                                                                                                                                                                                                                                                                                93US-00042747.
  3.2%;
                                                                        CCCTGCCACGGTGCACC 1
                                                                                                                                                                                                                                                                                                                                                                       93US-00042747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22 TGACCGAGGGCTGGGAC
                                                                                                                                              AAT16477 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                (first entry)
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15; Conservative
  Query Match
Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT32058 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1996-105220/11.
                                                                                                                                                                                                                            Sense primer
                                                                                                                                                                                                                                                                                                                                                                                              01-APR-1993;
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                                                                                                                                                                                                                                                                                                                                              30-JAN-1996.
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                                                                                                                                                                                                                                                                                            Synthetic.
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Best Local S
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                                                                                                                                                                          AAT16477;
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Wed Apr 21 12:58:21 2004
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The present sequence is an antisense oligomuclectide specific for the HIV target site, tat, which can be used for the detection of HIV, or for the treatment of HIV infection. The oligomuclectide has an average EC(90) (IM) of 1500, which refers to the conc. of oligomuclectide required to achieve 90 % inhibition of HIV p24 core antigen prodn. (contg. phosphorothioate linkages only)
                                                                                                                                                                                                                                                                                       Oligo:nucleotide(s) corresponding to HIV sequences - used for the detection of HIV or for inhibiting HIV propagation, partic. in infected
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human immunodeficiency virus; HIV; antisense oligonucleotide; tat;
detection; treatment; infection; inhibition; p24; core antigen;
production; ss.
                       Human immunodeficiency virus; HIV; antisense oligonucleotide; tat; detection; treatment; infection; inhibition; p24; core antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    / Match 3.2%; Score 13.8; DB 1; Length 21; Local Similarity 88.2%; Pred. No. 3.2e+02; nes 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 2 A; 7 C; 6 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HIV tat targetting antisense oligonucleotide.
tat targetting antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                              Example 3; Page 50; 90pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              240 gacracrircccagacric 256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAT32083 standard; RNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       94US-00277857.
                                                                                                                                                           95WO-US009080.
                                                                                                                                                                                   94US-00277857
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GGATGCTTCCAGGGCTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GENP-) GEN-PROBE INC
                                                                                                                                                                                                            (GENP-) GEN-PROBE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kwoh TJ;
                                                                                                                                                                                                                                      Ryder TB, Kwoh TJ;
                                                                                                                                                                                                                                                                WPI; 1996-105849/11
                                                     production; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-JUL-1995;
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                                                                                                                                                                                     19-JUL-1994;
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                                                                                                                                                           14-JUL-1995;
                                                                                                       WO9602557-A1
                                                                                                                                  01-FEB-1996,
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                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAT32083;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 211
  HIV
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The present sequence is an oligomucleotide complementary to an antisense oligomucleotide specific for the HIV target site, tat, which can be used for the detection of HIV, or for the treatment of HIV infection. The DNA equivalent of the antisense oligomucleotide has an average EC(90) (mN) of 1500, which refers to the conc. of oligomucleotide required to achieve 90 % inhibition of HIV p24 core antigen prodn. (contg. phosphorothicate
                                                                                                                                    The present sequence is an antisense oligonucleotide specific for the HIV target site, tat, which can be used for the detection of HIV, or for the treatment of HIV infection. The DNA equivalent of the oligonucleotide has an average EC(90) (nM) of 1500, which refers to the conc. of oligonucleotide required to achieve 90 % inhibition of HIV p24 core antigen prodn. (contg. phosphorothicate linkages only)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligo:nucleotide(s) corresponding to HIV sequences - used for the detection of HIV or for inhibiting HIV propagation, partic. in infected
                                         Oligo:nucleotide(s) corresponding to HIV sequences - used for the detection of HIV or for inhibiting HIV propagation, partic. in infected
                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Oligonucleotide complementary to HIV tat targetting antisense oligo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human immunodeficiency virus; HIV; antisense oligonucleotide; tedetection; treatment; infection; inhibition; p24; core antigen; production; complementary; ss.
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                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 64.7%; Pred. No. 3.2e+02;
Matches 11; Conservative 4; Mismatches 2; Indels
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                                                                                                                                                                                                                                                      Sequence 21 BP; 2 A; 7 C; 6 G; 0 T; 6 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 3; Page 69; 90pp; English.
                                                                                                           Example 3; Page 56; 90pp; English
                                                                                                                                                                                                                                                                                                                                                         240 GGCTGCTTCCCGGGCTC 256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP.
                                                                                                                                                                                                                                                                                                                                                                              || :||:|| |||:|
|GAUGCUUCCAGGGCUC 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   94US-00277857.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             134/c
AAT32134 standard; RNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1996-105849/11.
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               WPI; 1996-105849/11.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAT32134;
                                                                               subjects.
                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 212
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Gaps

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AAD19719;
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                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence is an oligonucleotide complementary to an antisense oligonucleotide specific for the HIV target site, tat, which can be used for the detection of HIV, or for the treatment of HIV infection. The antisense oligonucleotide has an average EC(90) (mM) of 1500, which refers to the conc. of oligonucleotide required to achieve 90 % inhibition of HIV p24 core antigen prodn. (contg. phosphorothicate
                                                                                                                                                                                                                                                                                                                                                                         corresponding to HIV sequences - used for the for inhibiting HIV propagation, partic. in infected
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                    Gaps
                                                                                                                                                             Oligonucleotide complementary to HIV tat targetting antisense oligo.
                                                                                                                                                                                immunodeficiency virus; HIV; antisense oligonucleotide; tat;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                    ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 13.8; DB 1; Length 21;
Pred. No. 3.2e+02;
0; Mismatches 2; Indels
3.2%; Score 13.8; DB 1; Length 21; 88.2%; Pred. No. 3.2e+02; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21 BP; 6 A; 6 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Simian herpesvirus B DNA primer B4.
                                                                                                                                                                                                                                                                                                                                                                                                             Example 3; Page 63; 90pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  240 GGCTGCTTCCCGGGCTC 256
                                     240 GGCTGCTTCCCGGGCTC 256
                                                                                                                                                                                                   production; complementary; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3.2%;
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                                                                                                                                                                                                                                                                                              94US-00277857
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA; 21
                                                        GGATGCTTCCAGGGCTC
                                                                                                     AAT32109 standard; DNA; 21
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                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 88.2
Matches 15; Conservative
          Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                (GENP-) GEN-PROBE INC.
                                                                                                                                                                                                                                                                                                                                                                        Oligo:nucleotide(s)
detection of HIV or
                                                                                                                                                                                                                                                                                                                                    Ryder TB, Kwoh TJ;
                                                                                                                                                                                                                                                                                                                                                      WPI; 1996-105849/11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAV33173 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linkages only)
                                                                                                                                                                                                                                                                                               19-JUL-1994;
                                                                                                                                                                                                                                                                            14-JUL-1995;
                                                                                                                                           16-SEP-1996
                                                                                                                                                                                                                                       WO9602557-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-NOV-1998
                                                                                                                                                                                                                                                          01-FEB-1996.
                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                            subjects.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAV33173;
                                                                                                                         AAT32109;
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                                                       17
  Query Match
                                                                                    RESULT 213
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                                                                                              AAT32109,
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The invention provides the Simian herpesvirus B DNA (AAV33167) sequence coding for a gB glycoprotein (UL27; AAW70293) and a portion of an ICP 15. Kba protein (UL28; AAW70294). The invention uses these DNA and protein sequences as a basis for the development of differential diagnostic tests for the rapid identification of Simian herpesvirus B cases. Primer BV1 (AAV33169) and BV2 (AAV33169), along with the Simian herpesvirus B sequence specific PBs probe (AAV33179), were used in these diagnostic tests. Other primer sets used were the sense primers B3 (AAV33171) or B3' (AAV33172) and antisense primers B4 or B4' (AAV33174). Therefore, the virus can be detected by detecting the DNA sequence and knowledge of the amino acid sequence will help in the design of DNA production
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Monkey herpes B virus DNA - coding for gB glycoproteins and polypeptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human, Mammary Gland Cancer Specific Gene, MSG, cytostatic, vaccine; cancer, therapy, immune response; PCR primer; ss.
Simian herpesvirus B gB glycoprotein; UL27; ICP protein; UL28; differential diagnostic test; immunoassay; antibody; PCR; primer; amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human MSG agmam023 cDNA amplifying agmam023 reverse PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.2%; Score 13.8; DB 1; Length 21; 88.2%; Pred. No. 3.2e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                Ä
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 2 A; 5 C; 9 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                Black
                                                                                                                                                                                                                                                                                                                                                                                                                Eberle R,
                                                                                                                                                                                                                                                                                                                                                               (SWBI-) SOUTHWEST FOUND BIOMEDICAL RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            English
                                                                                                                                                                                                                                                                                                                                                                                                                Hilliard J, Scinicariello F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAD19719 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22 TGACCGAGGCTGGGAC 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-MAR-2000; 2000US-0192277P.
                                                                                                Synthetic.
Cercopithecine herpesvirus 1.
                                                                                                                                                                                                                                                                   95US-00541878
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-MAR-2001; 2001WO-US009525
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 8; Col 7-8; 22pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.22
Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1998-361791/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200172780-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                   10-OCT-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18-DEC-2001
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                                                                                                                                                                        US5767265-A.
                                                                                                                                                                                                                      16-JUN-1998.
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Wed Apr 21 12:58:21 2004

Hu P, Recipon H, Cafferkey R;

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Sequence 21 BP; 3 A; 2 C; 9 G; 7 T; 0 U; 0 Other;
                                                     Example 3; Page 77; 99pp; English
                                                                                                                                                                                                                    AAH89013 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                                                                                                               10-NOV-1999; 99US-0164596P.
                                                                                                                                                                                                                                                                                                                                      10-NOV-2000; 2000WO-US030766
                                                                                                                                                                                                                                        27-FEB-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                          (GLAX ) GLAXO GROUP LTD. (AFFY-) AFFYMETRIX INC.
                                                                                                                                           tissues and organisms
    (DIAD-) DIADEXUS INC.
                       WPI; 2001-616468/71.
                                                                                                                                                                                                                                                                                                                   WO200134840-A2
                                            gland cancer.
                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                            17-MAY-2001.
              Salceda S,
                                                                                                                                                                                                                                                                                           Key
Variation
                                                                                                                                                                                                                              AAH89013;
                                                                                                                                                                                                           RESULT 216
                                                                                                                                                                                                                AAH89013,
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85 CAGTGGACATCACCACG 101

21 CACTAGACATCACCACG S

Location/Qualifiers

replace(11,a)

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The present invention describes the use of irinotecan (I) or its derivative for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a cytochrome p450, subfamily IIIA (nifedipine cytostatic applied (II). (I) and (II) have cytostatic activity. The therapeutic applications of (I) is improved, since it is possible to individually treat a subject with an appropriate
                                                                                                                                                                                                                           The present invention relates to human oligonucleotides comprising a single nucleotide polymorphic site (SNP: AAH88797-AAH89219). The present sequence is one such oligonucleotide. The oligonucleotides can be used in forensics, paternity testing, correlation of polymorphisms with phenotypic traits, genetic mapping of phenotypic traits and marker assisted breeding of animals and crop plants
                                                                                                           disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New use of irinotecan for preparation of compositions for treating in subject having genome with variant allele comprising cytochrome subfamily IIIA, polypeptide 5 polymucleotide, termed CYP3A5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cancer; CYP3A5; irinotecan; pharmaceutical; malignant glioma; cytochrome p450; subfamily IIIA; nifedipine oxidase; polypeptide cytostatic; PCR primer; 8s.
                                                                                 New polymorphic sites derived from the human genome are useful to determine sites correlating with phenotypic traits, particularly and also in forensics and paternity testing.
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Match 3.2%; Score 13.8; DB 1; Length 21; Local Similarity 88.2%; Pred. No. 3.2e+02; hes 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cancer based on CYP3A5 related oligonucleotide SEQ ID NO:4.
                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 2 A; 8 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
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Thomas
                                                                                                                                                                                    Claim 68; Page 11; 43pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    75 GAGGGCCGCGCAGTGGA 91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-JUL-2002; 2002WO-EP008219.
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24-MAY-2002; 2002EP-00011710.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACF62203 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-OCT-2003 (first entry)
Patil N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-268144/26.
                                           WPI; 2001-335945/35.
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Chen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Heinrich G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ACF62203;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               셤
                                                                                                                                                                                                                                                                                                    The present sequence is a PCR primer used for amplifying human mammary gland cancer specific gene (MSG) CDNA. MSG is useful for diagnosing, detecting, monitoring, treaging, incaping and treating mammary gland cancer in a patient by determining the levels of MSG in cells, tissues or bodily fluids in a patient and comparing the determined levels of MSG with levels of MSG in cells, tissues or bodily fluids from a normal human control, where a change in determined levels of MSG in the patient versus normal control is associated with the presence of mammary gland cancer. MSG is used for identifying potential therapeutic agents for use in imaging and treating mammary gland cancer. MSG and treating mammary gland cancer in a patient. MSG vaccine is useful for treating mammary gland cancer in a patient. MSG vaccine is useful for inducing an immune response against a MSG protein and for treating mammary gland cancer in a patient. MSG and its protein and for treating mammary gland cancer in a patient. MSG and its protein and for treating mammary gland cancer and for diagnosis and treatment of disorders of cells,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                   New isolated polynucleotide, mammary gland cancer specific gene (MSG), useful for diagnosing, monitoring, staging, imaging and treating mammary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; single nucleotide polymorphic; SNP; forensic science;
paternity testing; phenotypic trait; genetic mapping; animal breeding;
plant breeding; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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/standard_name= "single nucleotide polymorphism"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 3.2%; Score 13.8; DB 1; Length 21; Best Local Similarity 88.2%; Pred. No. 3.2e+02; Matches 15; Conservative 0; Mismatches 2; Indels
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dosage and/or an appropriate derivative of (1). Therefore, undesirable, harmful or toxic effects are efficiently avoided. Unnecessary and potentially harmful treatment of those subjects who do not respond to the treatment with substances (nonresponders), as well as the development of tway resistances due to suboptimal drug dosing can be avoided. ACF62200 to ACF62751 and ABM34912 to ABM35013 represent sequences used in the exemplification of the present invention
   88888888888
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Sequence 21 BP; 6 A; 8 C; 6 G; 0 T; 0 U; 1 Other;

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Gaps
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 3.2%; Score 13.8; DB 1; Length 21; 78.9%; Pred. No. 3.2e+02; tive 1; Mismatches 3; Indels
                                                                       336 GACCAGGCCGGCTGCTCT 354
                                                                                                        21 GTCCTGGGCCKGCTGCTGT 3
Query Match 3.2'
Best Local Similarity 78.9
Matches 15; Conservative
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Cancer; CYP3A5; irinotecan; pharmaceutical; malignant glioma; cytochrome p450; subfamily IIIA; nifedipine oxidase; polypeptide 5; cytostatic; PCR primer; ss. Cancer based on CYP3A5 related oligonucleotide SEQ ID NO:3. ACF62202 standard; DNA; 21 BP (first entry) WO2003013534-A2. 08-OCT-2003 Synthetic. ACF62202; RESULT 218 ACF62202 

23-JJL-2002; 2002WO-EP008219. 20-FEB-2003.

(EPID-) EPIDAUROS BIOTECHNOLOGIE AG. 23-JUL-2001; 2001EP-00117608. 24-MAY-2002; 2002EP-00011710.

Heinrich G, Kerb R;

WPI; 2003-268144/26.

New use of irinotecan for preparation of compositions for treating cancer in subject having genome with variant allele comprising cytochrome p450, subfamily IIIA, polypeptide 5 polynucleotide, termed CYP3A5.

Disclosure; Page 32; 86pp; English.

The present invention describes the use of irinotecan (I) or its derivative for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a cytochrome p450, subfamily IIIA (nifedipine cytostatic activity. The therapeutic applications of (I) is improved, since it is possible to individually treat a subject with an appropriate cytostatic activity. The therapeutic applications of (I) is improved, since it is possible to individually treat a subject with an appropriate derivative of (I). Therefore, undesirable, harmful or toxic effects are efficiently avoided. Unnecessary and potentially harmful treatment of those subjects who do not respond to the treatment with substances (nonresponders), as well as the development of the treatment with substances (nonresponders), as well as the development of charge resistances and ABM34912 to ABM36013 represent sequences used in the exemplification of the present invention

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Gaps ö

3.2%; Score 13.8; DB 1; Length 21; 78.9%; Pred. No. 3.2e+02; tive 1; Mismatches 3; Indels

336 GACCAGGGCCGGCTGCTCT 354

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Local Similarity 78.9 les 15; Conservative

Matches

Query Match

21 GTCCTGGGCCKGCTGCTGT 3

RESULT 220

Sequence 21 BP; 0 A; 6 C; 8 G; 6 T; 0 U; 1 Other;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes a method for the use of irinotecan (I) or its derivative for the preparation of a pharmaceutical composition for treating colorectal, crivical, gastric, lung, ovarian or pancreatic cancer, or malignant gloma in a subject having a genome with a variant allele which comprises a multidrug resistance protein 1 (MRP1) allele which comprises a multidrug resistance protein 1 (MRP1) (I) or its derivative can be used for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject, where the subject is a human (preferably African or Asian) or a muouse. The present sequence represents a sequence which is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                               irinotecan; colorectal cancer; cervical cancer; gastric cancer;
lung cancer; ovarian cancer; pancreatic cancer; malignant glioma;
variant allele; multidrug resistance protein 1; MRP1; cytostatic; gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Use of irinotecan or its derivative for preparation of a pharmaceutical composition for treating cancer in a subject having a genome with a variant allele comprising a multidrug resistance protein 1
                                            Gape
                                          ö
        Length 21;
    3.2%; Score 13.8; DB 1; Length 2:
78.9%; Pred. No. 3.2e+02;
tive 1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                           MRP1 based cancer related nucleic acid SEQ ID NO:4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 6 A; 8 C; 6 G; 0 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 41; 100pp; English.
                                                                               336 GACCAGGCCGGCTGCTCT 354
                                                                                                                 1 Greeregecekeereer 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23-JUL-2001; 2001EP-00117608.
24-MAY-2002; 2002EP-00011710.
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                                                                                                                                                                                                               ADB20874 standard; DNA; 21
3.2%
Best Local Similarity 78.9%
Matches 15; Conservative
                                                                                                                                                                                                                                                                                      20-NOV-2003 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Heinrich G, Kerb R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO2003013533-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polynucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unidentified.
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ADB20873;

ADB20873

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The invention relates to the novel use of irinotecan to treat a patient suffering from cancer. This involves determining if the patient has one or more variant alleles of the UGTIAL gene, and if the patient has one or more of such variant alleles, irinotecan is administered in an increased or decreased amount in comparison to the amount that is administered without regard to the patient's alleles in the UGTIAL gene. The invention has cytostatic activity. A composition of the invention acts as a topoisomerase I inhibitor. The method is useful for treating a patient, an animal e.g. mouse or a human, preferably African or Asian, suffering from cancer such as colorectal, cervical, gastric cancer, lung, ovarian, pancraatic cancer or malignant glioma. The present sequence is udes in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Use of irinotecan to treat cancer patient by determining if patient has variant alleles of UGTIA1 gene, administering increased/decreased amounts of irinotecan based on increased/decreased levels of UGTIA1 gene product.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ss; irinotecan; cancer; UGTIA1; cytostatic; topoisomerase I inhibitor; colorectal cancer; cervical cancer; gastric cancer; lung cancer; ovarian cancer; pancreatic cancer; malignant glioma; uridine diphosphate glycosyltransferasel member Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               / Match 3.2%; Score 13.8; DB 1; Length 21; Local Similarity 78.9%; Pred. No. 3.2e+02; Nes 15; Conservative 1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human UGTIAl variant allele sequence fragment SEQ ID NO:3.
ovarian cancer; pancreatic cancer; malignant glioma; uridine diphosphate glycosyltransferasel member Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 6 A; 8 C; 6 G; 0 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                 (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 8; Page 44; 107pp; English
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                                                                                                                                                                                                                 23-JUL-2002; 2002WO-EP008217.
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24-MAY-2002; 2002EP-00011710.
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                                                                                                                                                                                                                                                                                                                                                                                 Heinrich G, Kerb R;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes a method for the use of irinotecan (1) or its derivative for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a multidrug resistance protein 1 (MRP1) golynucleotide (II). (I) has expostatic activity. (I) or its derivative can be used for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject, where the subject is a human (preferably African or Asian) or a mouse. The present sequence represents a sequence which is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Use of irinotecan or its derivative for preparation of a pharmaceutical composition for treating cancer in a subject having a genome with a variant allele comprising a multidrug resistance protein 1 polynucleotide.
                                                                                                                                                                                                                         irinotecan; colorectal cancer; cervical cancer; gastric cancer;
lung cancer; ovarian cancer; pancreatic cancer; malignant glioma;
variant allele; multidrug resistance protein 1; MRP1; cytostatic; gene;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human UCT1A1 variant allele sequence fragment SEQ ID NO:4.
                                                                                                                                                                            WRP1 based cancer related nucleic acid SEQ ID NO:3,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 41; 100pp; English.
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24-MAY-2002; 2002EP-00011710.
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                                          ADB20873 standard; DNA; 21
                                                                                                                                    (first entry)
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                   Unidentified.
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ADB87963;

Best Loca Matches

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Gaps

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The invention relates to the novel use of irinotecan to treat a patient suffering from cancer. This involves determining if the patient has one or more variant alleles of the UGTAH gene, and if the patient has one or more of such variant alleles, irinotecan is administered in an increased or decreased amount in comparison to the amount that is administered without regard to the patient's alleles in the UGTAH, gene. The invention has cytostatic activity. A composition of the invention acts as a topoisocmerase I inhibitor. The method is useful for treating a patient, an animal e.g. mouse or a human, preferably African or Asian, suffering from cancer such as colorectal, cervical, gastric cancer, lung, ovarian, pancreatic cancer or malignant glioma. The present sequence is udes in the exemplification of the invention.
                                                                                                                                                                                                                Use of irinotecan to treat cancer patient by determining if patient has variant alleles of UGTIA1 gene, administering increased/decreased amounts of irinotecan based on increased/decreased levels of UGTIA1 gene product.
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                                                                      (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
                                                                                                                                                                                                                                                                                                                      Claim 8; Page 44; 107pp; English
23-JUL-2001; 2001EP-00117608.
24-MAY-2002; 2002EP-00011710.
                                                                                                                       Heinrich G, Kerb R;
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3.2%; Score 13.8; DB 1; Length 21; 78.9%; Pred. No. 3.2e+02; ive 1; Mismatches 3; Indels Local Similarity 78.5 Query Match Best Loca Matches

336 GACCAGGCCGGCTGCTCT 354 1 GTCCTGGGCCKGCTGCTGT 19 ઠ 셤

ADB96945 standard; DNA; 21 ADB96945; RESULT 223 ADB96945 ##X#X#X#X#X#X#X#X#X#X#X#X#X#X#X#X#X#

(first entry) 04-DEC-2003

Human UGT1A1 variant allele sequence fragment SEQ ID NO:3.

irinoțecan; colorectal cancer; cervical cancer; gastric cancer; lung cancer; ovarian cancer; pancreatic cancer; malignant glioma; multidrug resistance 1; MDR1; cytostatic; human; ds; Cyp3A5; MRP1; MDR1;

Homo sapiens.

WO2003013537-A2.

20-FEB-2003.

23-JUL-2002; 2002WO-EP008218. 23-JUL-2001; 2001EP-00117608 24-MAY-2002; 2002EP-00011710 (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.

Heinrich G, Kerb R;

New use of irinotecan for preparation of pharmaceutical compositions for treating cancer in subject having genome with variant allele comprising

multidrug resistance 1 polymucleotide.

Disclosure; Page 69; 130pp; English.

The invention relates to the novel use of irinotecan or its derivative for the preparation of pharmaceutical compositions for treating colorectal, cervical, gateric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a multidrug resistance I (MDR1) polymucleotide. A composition of the invention has cytostatic activity. The invention is useful for the preparation of pharmaceutical compositions for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject (preferably human, more preferably African or Asian) or a mouse. The present sequence is used in the exemplification of the invention. 

Sequence 21 BP; 0 A; 6 C; 8 G; 6 T; 0 U; 1 Other;

Gaps .. 0 Score 13.8; DB 1; Length 21; Pred. No. 3.2e+02; 1; Mismatches 3; Indels 3.2%; 15; Conservative Local Similarity Query Match Matches

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ADB96946 standard; DNA; 21 RESULT 224

ADB96946;

(first entry)

04-DEC-2003

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Human UGT1A1 variant allele sequence fragment SEQ ID NO:4.

irinotecan; colorectal cancer; cervical cancer; gastric cancer; lung cancer; ovarian cancer; pancreatic cancer; malignant glioma; multidrug resistance 1; MDR1; cytostatic; human; ds; Cyp3A5; MRP1; MDR1; TOP1.

Homo sapiens.

WO2003013537-A2.

20-FEB-2003.

23-JUL-2002; 2002WO-EP008218.

23-JUL-2001; 2001EP-00117608. 24-MAY-2002; 2002EP-00011710.

(EPID-) EPIDAUROS BIOTECHNOLOGIE AG.

я, Heinrich G,

WPI; 2003-268145/26.

New use of irinotecan for preparation of pharmaceutical compositions for treating cancer in subject having genome with variant allele comprising multidrug resistance 1 polynucleotide.

Disclosure, Page 69, 130pp; English.

The invention relates to the novel use of irinotecan or its derivative for the preparation of pharmaceutical compositions for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant gliona in a subject having a genome with a variant allele which comprises a multidrug resistance 1 (MDR1) polynucleotide. A composition of the invention has cytostatic activity. The invention is useful for the preparation of pharmaceutical compositions for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant ADB96946/6/MD96946/4/MD96946/4/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD96946/MD9694/MD9694/MD9694/MD9694/MD9694/MD9694/MD9694/MD9694/MD9694/MD9694/MD9694/MD9694/MD9694/MD96

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ADB92137, RESULT

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The invention relates ro a novel use of irinotecan or its derivative for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant allel and subject having a genome with a variant allele which comprises a multidrug resistance 1 (MRM1) polynucleotide. A composition of the invention has cytostatic activity. The present sequence is used in the exemplification of the invention.
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                                                                                                                                                             irinotecan; colorectal cancer; cervical cancer; gastric cancer;
lung cancer; ovarian cancer; pancreatic cancer; malignant glioma;
multidrug resistance 1; MDR1; cytostatic; ds; human; UGT1A1; MRP1; TOP1.
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                                                                                                                         Human UGT1A1 variant allele sequence fragment SEQ ID NO:3.
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24-MAY-2002; 2002EP-00011710.
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  ADB92136 standard; DNA; 21
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Best Local Similarity 78.9'
Matches 15; Conservative
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10-MAR-2003
20-APR-1993
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glioma in a subject (preferably human, more preferably African or Asian) or a mouse. The present sequence is used in the exemplification of the
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                                                                                                                         Length 21;
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                                                                                                                    Query Match
3.2%; Score 13.8; DB 1; Length 2
Best Local Similarity 78.9%; Pred. No. 3.2e+02;
Matches 15; Conservative 1; Mismatches 3; Indels
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24-MAY-2002; 2002EP-00011710
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                                        invention.
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Gaps

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Feighner SD;

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Comparative analysis of regions close to both the 3' and 5' ends of small subunit ribosomal RNA sequences with near identity in the enkaryotic kingdom identified two consensus sequences, ERIB 1 and ERIB 10, spanning the serRNA gene sequence. These primers may be used in PCR to selectively mapped from a multiply the serRNA genes contained within the genomic DNA prepach from a number of Einmeria species, to determine the degree of similarity between serRNA from different Einmeria species. The probe CommonARC represents a sequence common to all Einmeria species which may be used to identify Einmeria infection. See also AAQ3128-332. NOVE: As specifications EP-516381. EP-516383-6, EP-516381 and EP-516395-6 are identical except in the claims section, sequences for all these specifications can be found indexed under EP-516385. However the claims dequences of each specification will be indexed under their own patent number, thus each specification will be indexed under their own patent number, thus each missing OS field,) (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PI field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Glutaminase, antiviral, virucide, anticancer; cancer therapy, HIV virus, gene therapy, Escherichia coli, primer; ss.
                                                                                                                                                                                                                   Species-specific Eimeria tenella DNA probes - comprise divergent DNA sequences and are complementary to E. tenella small sub-unit ribosomal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3.2%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 3.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 4 A; 4 C; 5 G; 7 T; 0 U; 0 Other;
                                                                                                                                  Elbrecht A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pseudomonas glutaminase primer JR-1.
                                                                                                                                                                                                                                                                                     Disclosure; Page 21; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             399 AAGGTCTTCTACGTGATCGA 418
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                                                                                                                                Chakraborty PR,
P-JuchelkaH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ68439 standard; DNA; 20 BP.
                92EP-00304781
                                                  91US-00707362.
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(first entry)
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                                                                                              (MERI ) MERCK & CO INC.
                                                                                                                                                                                   WPI; 1992-400736/49.
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                                                                                                                                Dashkevicz M,
                                                                                                                                                    Liberator PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9413817-A1
                27-MAY-1992;
                                                29-MAY-1991;
                                                                12-MAY-1992;
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12-JAN-1995
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                                                                                                                                                                                                                                                     RNA gene
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Matches
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Oligo-nucleotide(s) able to inhibit multi:drug resistant phenotype - either by anti:sense or aptameric effects, useful for enhancing cytotoxic effects of chemotherapeutic agents on multi:drug resistant cancer cells.
                                                                                                       Recombinant glutaminase derived from Pseudomonas 7A - expressed in E. coli to increase yield and avoid Pseudomonas endotoxins for antiviral and anticancer therapy.
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                                                                                                                                                                                                                                                                   Chromosomal DNA from Pseudomonas sp. 7A (ATCC 29598) was used to construct a genomic library in Escherichia coli LE392. Screening with mixed oligomucleotide probes was used to isolate a gluteminase-encoding clone. This was sequenced using the primers given in AAQ68439-47. The gene can be used to manufacture recombinant glutaminase, free of Pseudomonas exotoxin, for use in e.g. HIV and cancer therapy. The gene may also be used in gene therapy protocols. (Updated on 25-MAR-2003 to correct PN field.)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human multidrug resistance-1; MDR-1; inhibition; aptameric; human multidrug resistance-associated protein; antisense; cytotoxic;
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Freeman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Complementary human MRP oligonucleotide OL(8E)MRP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3.2%; Score 13.6; DB 1;
80.0%; Pred. No. 3.2e+02;
ative 0; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 3 A; 5 C; 7 G; 5 T; 0 U; 0 Other;
Sethuraman N,
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                                                                                                                                                                                                                           Disclosure, Fig 2B; 60pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     265 TGCACCTGGAGCAGGGCGGC 284
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Roberts J, Macallister TW,
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                                                      WPI; 1994-217891/26.
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misc_feature
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Matches
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92WO-US010421 92WO-US010421

04-DEC-1992; 04-DEC-1992;

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The polypeptides of the invention may be of use in treating these
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Vaccine, eye disease, conventional trachoma; nonendemic trachoma; paratrachoma; inclusion conjunctivitis; genital disease, perihepatitis; nongonococcal uretritis; epidymitis; cervicitis; salpingitis; PCR primer; bartholinitis; pneumopathy; venereal lymphogranulomatosis; ss.
                            The present sequence represents a novel oligonucleotide OL(8E)MRP that specifically hybridises in a human cell with a complementary sequence of human multidrug resistance-associated protein (MRP) gene. Hybridisation causes inhibition of expression of the multidrug resistance phenotype by the cell, due to the oligonucleotide having an aptementic inhibitory effect as well as an antisense inhibitory effect. The oligonucleotide definition of assertant phenotype. When co- administered to cancer patients to prevent development of the multidrug resistant phenotype. When co- administered with Chemotherapeutic agents, the oligonucleotide is useful for potentiating elimination of multidrug resistant tumour cells from bone marrow or peripheral stem cell grafts.
                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                             Also, the oligonucleotide can be used as an immunosuppressive agent
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                                                                                                                                                                                                                                                     3.2%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 3.2e+02; Live 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR primer used to amplify an ORP of Chlamydia trachomatis.
                                                                                                                                                                                                                          Sequence 20 BP; 4 A; 2 C; 11 G; 3 T; 0 U; 0 Other;
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   Disclosure; Page 17; 74pp; English.
                                                                                                                                                                                                                                                                                                                       28 AGGGCTGGGACGAAGATGGC 47
                                                                                                                                                                                                                                                                                                                                             1 AGGGCGGGATGATGGC 20
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97FR-00016034.
98US-0107077P.
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                                                                                                                                                                                                                                                                                        16; Conservative
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17-DEC-1997;
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Vaccine; eye disease; conventional trachoma; nonendemic trachoma; paratrachoma; inclusion conjunctivitis; genital disease; perihepatitis; nongonococcal uretritis; epidymitis; cervicitis; salpingitis; PCR primer; bartholinitis; pneumopathy; venereal lymphogranulomatosis; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PCR primer used to amplify an ORF of Chlamydia trachomatis.
                                                    20;
                                                                                                           4; Indels
                                                    Length
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Sequence 20 BP, 3 A, 4 C, 6 G; 7 T; 0 U; 0 Other;
                                                    Score 13.6; DB 1;
Pred. No. 3.2e+02;
                                                                                                              0; Mismatches
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                                                                                                                                                                      120 AAGTACGGCATGCTGGCCCG 139
                                                                                                                                                                                                        20 AAATACGCCATGCTGACCAG 1
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                                                       Query Match
Best Local Similarity 80.0%;
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                                                                                                                  16, Conservative
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17-DEC-1997;
04-NOV-1998;
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                                                                                                                                        Respiratory disease; pneumonia; bronchitis; heart disease; sarcoidosis; sinusitis; purulent otitis media; erythema nodosum; pharyngitis; vaccine; neutralising epitope; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human leukocyte antigen; HLA; class I allele type; probe; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3.2%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 3.2e+02; ive 0; Mismatches 4; Indels
                                                                                                              PCR primer used to amplify an ORF of Chlamydia pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 5 A; 6 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Genome sequence of Chlamydia pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Page 1724; Disclosure; 1912pp; English.
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                                                                                                                                                                                                                                                                                                                                   97FR-00014673
98US-0107078P
                           AAX95138 standard; DNA; 20
                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    especially where the vect epitope of C. pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16; Conservative
                                                                                                                                                                                                                 Chlamydophila pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1999-357842/30.
                                                                                                                                                                                                                                                                                                                                                                            GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                   21-NOV-1997;
04-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-OCT-2000
                                                                                                                                                                                                                                                                                                      20-NOV-1998;
                                                                                                                                                                                                                                              WO9927105-A2
                                                                                    13-SEP-1999
                                                                                                                                                                                                                                                                            03-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                      Griffais R;
                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAA67067;
                                                        AAX95138;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 233
RESULT 232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAA67067
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The present invention describes a method for distinguishing a human betworpte antigen of alse I antigen or allele by a combination of polymerase chain reaction (PCR) using a primer pair whereby all HLA-A, -B or -C alleles can be amplified or using reverse hybridisation analysis comprising a DNA probe covalently bonded to microtire plate wells which are hybridisable specifically with the base sequence of at least one specific HLA-A, -B or -C allele. The method is applicable in gene typing, judying donor-recipient compatibility during organ transplant and correlation analysis for diagnosis of various diseases. The method is simple, rapid and accurate, with possibility of mechanisation and techniques. AAA666943 to AAA67072 represent oligonucleotide probes and PCR primers for use in the method of the present invention
                                                                                                                                                                                                                                                                                                           Simple, rapid and accurate method for distinguishing HLA class I allele type with possibility of mechanization and automation, applicable in judging donor-recipient compatibility during organ transplant and disease
amplification; hybridisation; organ transplant; gene typing; diagnosis;
ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4; Indels , 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           expression library; antibody; immunization; anchor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3.2%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 3.2e+02; tive 0; Mismatches 4; Indels ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Primer F3c used to amplify part of llama antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 3 A; & C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          187 CACATATCCACTGCTCGGTG 206
                                                                                                                                                                                                                                                                                                                                                                                           Claim 8; Page 78; 83pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20 cacacarccacresasedere 1
                                                                                                                                                                                     98JP-00335151.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-JAN-2000; 2000WO-EP000296.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAA73749/c
ID AAA73749 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                     (SHIO ) SHIONOGI & CO LID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 80.0
Matches 16; Conservative
                                                                                                                                                                                                                                                   Kaneshige T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (revised)
                                                                                                                                                                                                                                                                                WPI; 2000-400097/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200043507-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Llama; primer;
framework; ss.
                                                                                           WO200031295-A1.
                                                               Homo sapiens.
                                                                                                                                                         07-OCT-1999;
                                                                                                                                                                                     26-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-SEP-2003
14-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27-JUL-2000.
                                                                                                                         02-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lama glama.
                                                                                                                                                                                                                                                   Moribe T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA73749;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 234
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ò
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Sequence 20 BP; 6 A; 3 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200183749-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08-NOV-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAS97449;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mus sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 236
AAS97449/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 *****************
                                                                                                                                                                                                                                                                                                The present invention relates to an expression library comprising synthetic or semi-synthetic mucleic acid sequences, not cloned from an immunized source, where the nucleic acid sequences are derived from mucagenised immunoglobulins that are naturally devoid of light chains. The library is useful for the preparation of antibodies having binding specificity for a target antigen which avoids the need for a donor to have been previously immunized with the target antigen. The recombination of heavy and light chains is avoided, therefore preventing the formation of molecules that are non-functional. The number of hypervariable respectoire of possible binding variants to be obtained. The present sequence is a PCR primer targeted to anchor regions in Ilama antibodies. The primers (AAA73745 to AAA73745) amplified the framework regions in the primers (AAA73745 to AAA73745 amplified the framework regions F1, F2, F2c, F3 and F4. (Updated on 15-SEP-2003 to standardise OS field)
                                                                                                                                                                                       Expression library comprising nucleic acids not cloned from an immunized source, derived from immunoglobulins naturally devoid of light chains, use for producing antibodies specific for a target antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human angiotensinogen gene; AGT; insulin-dependent diabetes mellitus; type 1 diabetes; dhromsome 1462-43; single nucleotide polymorphism; IDDM; SNP; diagnosis; susceptibility; transgenic animal; drug screening; antidiabetic; gene therapy; alternative exon 1; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human angiotensinogen gene alternative exon 1 PCR primer, SEQ ID NO:17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 3.2%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 3.2e+02; Matches 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 2 A; 8 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Shattuck DM;
                                                                                                                                                                                                                                                                  Example 2; Page 29; 60pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       269 CCTGGAGGGGGGCACCA 288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CCTGGGGCCTGGCGGAACCA 1
                                                                                                            Van Der Logt CPE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAC91615 standard; DNA; 20 BP
99EP-00300351.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16-MAY-2000; 2000WO-US013327.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-MAY-1999; 99US-0135423P.
06-JAN-2000; 2000US-0174700P.
                                                                        (HIND-) HINDUSTAN LEVER LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (MYRI-) MYRIAD GENETICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mcgrail M, Russell DL,
                                    (UNIL ) UNILEVER PLC. (UNIL ) UNILEVER NV.
                                                                                                                                                   WPI; 2000-482910/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-025172/03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200071751-A1.
                                                                                                              Frenken LGJ,
19-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-NOV-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAC91615;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20
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AAC91615/c
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The invention relates to the human angiotensingen (AGT) gene, some mutant alleles of which cause a susceptibility to insulin-dependent diabetes of which cause a susceptibility to insulin-dependent confabetes mellitus (IDDM, type I diabetes). The AGT gene is located on chromosome 1442-43, a region linked to IDDM. The invention discloses genomic sequences comprising an alternative AGT gene (AAC91600-CC G91604) and a genomic sequence comprising an alternative AGT gene (AAC91600). The invention also encompasses the specifically claimed human AGT mutant nucleic acid sequences AAC91667-C91684, and the wutant confection mutant AGT alleles or gene products thereof which are related to detecting mutant AGT alleles or gene products thereof which are related to detecting mutant AGT alleles or gene products thereof which are related to confection of a polymorphism in the AGT gene; and methods of screening for drug candidates which may be useful in the treatment of diabetes resulting from an AGT mutation. Methods of preventing or treating diabetes are claimed which comprise the administration of a compound which agonises or antagonises wild-type or mutant AGT, which agonises or antagonises wild-type or mutant AGT, which agonises a transgenic non-human animal, or cell line derived compasses a transgenic non-human animal, or cell line derived compasses a transgenic non-human animal, or cell line derived compasses a transgenic non-human animal, or cell line derived care useful for secreening compounds which bind to AGT correcting or fragments thereof are useful for secreening compounds which bind to AGT correcting or fragments thereof are useful for secreening compounds which bind to AGT proteins or fragments thereof are useful for secreening compounds which bind to AGT proteins or fragments thereof are useful for secreening compounds which bind to AGT proteins and the pagence are useful for secreening of the invention of the invention
Novel angiotensinogen gene, mutant alleles of which causes susceptibility to insulin-dependent diabetes mellitus useful for diagnosis of predisposition to diabetes.
                                                                                                                                                                                                                                                                                    Example 2; Page 33; 83pp; English.
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. 0 Query Match 3.2%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 3.2e+02; Matches 16; Conservative 0; Mismatches 4; Indels

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AAS97449 standard; DNA; 20 BP. (first entry) Murine SAC1 gene-specific oligonucleotide PCR primer #54.

Human; mouse; SAC1; carbohydrate; sweetener; ethanol; alcoholism; ss; obesity; diabetes; transgenic embryo; body tissue; body fluid; pancreas; blood; tongue; PCR primer; anorectic; antidiabetic; gene therapy; protein replacement therapy;

25-APR-2001; 2001WO-US013387.

28-APR-2000; 2000US-0200794P. 28-JUL-2000; 2000US-0221419P. 10-NOV-2000; 2000US-0247443P.

(WARN ) WARNER LAMBERT CO.

(MONE-) MONELL CHEM SENSES CENT 

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Ξ
 De Jong PJ,
 Chatterjee A,
Tordoff MG;
 Bachmanov AA, Beauchamp GK,
Ohmen JD, Reed DR, Ross D,
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WPI; 2002-075162/10.

Novel isolated polypeptide comprising variant form of mouse or human SAC1 polypeptide, and is associated with altered preference for carbohydrates or other sweeteners, useful for preventing obesity, diabetes, alcoholism.

Claim 14; Page 75; 239pp; English.

The invention relates to an isolated polypeptide, comprising a variant form to f mouse or human SAC1 polypeptide. The variant form is associated with altered preference for carbohydrates, other sweeteners or ethanol. The polypeptide and its associated DNA sequence can be produced by recombinant techniques and is useful for preventing obssity, diabetes or alcoholism associated with SAC1 expression. The sequences are useful in creening for drugs and sweeteners. Recombinant cell lines and transgenic mbryos may be used in screening for and identifying agents that induce or repress function of SAC1. Predisposition to diabetes, obesity or alcoholism can be ascertained by testing any fluid or tissue of a human (such as blood, pancreas or tongue) for sequence variations of the SAC1 gene. A sequence variation of the SAC1 locus may indicate a predisposition to diabetes, obesity and/or alcoholism and may provide a diagnostic mark. The polymulectide can be detected in a biological sample by contacting the DNA with a probe to form a hybridisation complex which is then detected. The sequences represent cDNA encoding human and mouse SAC1 polypeptides and PCR primers specific for the SCA1 genes

Sequence 20 BP; 4 A; 8 C; 4 G; 4 T; 0 U; 0 Other;

Gaps ö Match 3.2%; Score 13.6; DB 1; Length 20; Local Similarity 80.0%; Pred. No. 3.2e+02; les 16; Conservative 0; Mismatches 4; Indels Query Match

7 GAGTGAAACTGCGGGTGACC 26

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20 GAGTGGAGCTGCAGGTTACC 1

ABL41764 standard; DNA; 20 BP. 237 ABL41764 RESULT

ABL41764;

29-MAY-2002 (first entry)

PCR primer used to amplify N-RAS proto-oncogene exon 2.

N-RAS; single base substitution; DNA mutation; PCR primer; ss.

Homo sapiens. Synthetic.

US6346386-B1.

12-FEB-2002.

29-SEP-2000; 2000US-00677045

29-SEP-2000; 2000US-00677045.

(ARUP-) ARUP INST.

Elenitoba-Johnson KSJ;

WPI; 2002-224990/28.

Determining mutation in DNA, comprises attaching guanine-cytosine-rich clamp to DNA, fluorescently labeling DNA and mixing it with denaturant,

heating to melt DNA and comparing melting temperatures of DNA and its wild type.

Example 3; Col 10; 16pp; English.

PCR primers ABL41762-64 were used to amplify exon 2 of the N-RAS protooncogene, in the course of the invention. The specification describes a
method for determining whether a DNA sequence contains an alteration. The
method comprises attaching a DNA sequent comprising one or more copies of
the DNA sequence to a guanine-cytosine-rich clamp, fluorescently labeling
the DNA sequence to a quanine-cytosine-rich clamp, fluorescently labeling
the DNA sequence to a guanine-cytosine-rich clamp, fluorescently labeling
and comparing the melting temperatures of the DNA segment and a wild type
alteration in the DNA sequence. The method is useful for determining
whether a DNA sequence contains an alteration. The method is suitable for
detecting a mutation as small as a single base substitution in a
relatively large DNA fragment. As the disparity in melting temperatures
is most evident in a lower melting domain of a DNA fragment, it is
possible to distinguish single base substitutions within lower melting domain

Sequence 20 BP; 9 A; 3 C; 6 G; 2 T; 0 U; 0 Other;

Gaps ô Query Match 3.2%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 3.2e+02; Matches 16; Conservative 0; Mismatches 4; Indels

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RESULT 238 ABQ74079

ABQ74079 standard; DNA; 20 BP.

ABQ74079;

(first entry) 11-OCT-2002 Microsatellite typing and sequencing D6S291 5' primer.

Homozygous stem cell; major histocompatibility complex; MHC; HLA; human leukocyte antigen; immunotype; genotype; microsatellite; probe; gent cell; nottorpic, neuroprotective; antigarkinsonian; vulnerary; cytostatic; antiatretriosclerotic; antiinflammatory; immunosuppressive; antianaemic; antidiabetic; tranquilliser; respiratory; cardiant; trauma; muscular; ophthalmological; gene therapy; genetic disease; cancer; cystic fibrosis; muscular dystrophy; cardiac condition; burn; myopathy; neurodegenerative disease; Alzheimer's disease; Parkinson's disease; miltiple sclerosis; post-trauma repair; reconstruction; blindness; disbetes; autoimmune disease; anaemia; PCR primer; ss.

Synthetic.

WO200257429-A2. 

25-JUL-2002.

02-JAN-2002; 2002WO-US000107.

02-JAN-2001; 2001US-0258881P.

(STEM-) STEMRON INC.

Yan WL;

WPI; 2002-575456/61.

Producing homozygous stem cells having a target genotype and/or immunotype from non-fertilized post-meiosis I diploid germ cells, suitable for diagnostic, therapeutic and cosmetic transplant and

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The present invention describes a method for producing homozygous stem (HS) cells having a target genotype and/or immunotype from non-fertilised post-meiosis I diploid germ cells by mitotically activating the germ cells to develop multiple blastocyst-like masses, each of which contains an inner cell mass (ICW) that is homozygous for the target genotype and/or immunotype. The methods of the present invention are useful for the production of HS cells utilised for diagnosis, therapeutic and cosmetic transplantation, cell replacement and/or gene therapy, and the treatment of various genetic diseases (Cystic fibrosis, muscular disease, Parkinson's disease and multiple sclerosis), traumatic injuries (post-traum repair and reconstruction, limb replacement, spinal cord injuries and burns), cancer, disorders of the epithelium (blindness, myopathy, atherosclerosis), Crohn's disease, autoimmune diseases and anaemia. ABQ74028 to ABQ74115 represent PCR primers and sequence specific oligonucleotide (SSO) probes which are used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             exemplification of the present invention
                                                   Disclosure, Fig 7; 75pp; English
treatment of various disorders.
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Sequence 20 BP; 3 A; 6 C; 7 G; 4 T; 0 U; 0 Other;

. 0 Score 13.6; DB 1; Length 20; Pred. No. 3.2e+02; 4; Indels 0; Mismatches 126 gecarecrescossories 145 1 GGCÁTTCAGGCATGCCTGGC 20 3.2%; 16; Conservative Local Similarity Query Match Best Loca Matches 셤 ð

ABZ88298 standard; DNA; 20 BP 17-OCT-2003 ABZ88298; RESULT 239 ABZ882 

(first entry)

Human oligonucleotide sequence.

Human, antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

sapiens OHOF 40200285308-A2.

31-OCT-2002

24-APR-2001; 2001US-0286137P.

23-APR-2002; 2002WO-US013135

(EPIG-) EPIGENESIS PHARM INC.

Aguilar D; Katz E, Pabalan J, Sandrasagra A, Ka ,, Shahabuddin S; Li Y, San Tang L, Nyce JW, Miller S,

WPI; 2003-229219/22.

Pharmaceutical composition for treating allments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Disclosure; SEQ ID NO 3540; 872pp; English

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the intiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an contintlammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallargic, antiasthmatic, hypotensive, contintiallargic, antiasthmatic, hypotensive, confortion is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antisinal ammatory steroid in a subject, for reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine copy, producing bronchodilation, increasing levels of ubiquinone or lung inflammation, lung allergies, or a respiratory disease or condition. Or specification, but was obtained in electronic format directly from WIPO continual products are septimated in the printed at the printed and the printed and the sequences.

Gaps ö Query Match
3.2%; Score 13.6; DB 1; Length 20;
Best Local Similarity 80.0%; Pred. No. 3.2e+02;
Matches 16; Conservative 0; Mismatches 4; Indels

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Gaps

ABZ92729 standard; DNA; 20 ABZ92729; ABZ92729 ID ABZ9 

BP

Human oligonucleotide sequence.

Human, antisense; lung dysfunction; nasal airway dysfunction; antiaflarmatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; dyrostatic; igene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

31-OCT-2002.

24-APR-2001; 2001US-0286137P. 23-APR-2002; 2002WO-US013135.

(EPIG-) EPIGENESIS PHARM INC.

Li Y, Sandrasagra A, Katz E, Pabalan J, Aguilar D; Tang L, Shahabuddin S;

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 7971; 872pp; English

¥888888888888888888888888

Sequence 20 BP; 5 A; 7 C; 5 G; 3 T; 0 U; 0 Other;

RESULT 240

(first entry) 17-OCT-2003

WO200285308-A2.

Nyce JW, L Miller S,

first active agent comprising an oligomorteotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or ergions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entiinflammatory steroid and ubiquinon. A composition of the invention has antiinflammatory, antiallargic, antiathmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antificial ammatory steroid in a subject, for reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition, Note: The sequence data for this patent is not represented in the printed appears in the printed of a perficult of the printed of the file of this patent is not represented in the printed of the file of this patent is not represented in the printed of the file of this patent is not represented in the printed of the file of this patent is not represented in the printed of the file of this patent is not represented in the printed of this patent is not represented in the printed of the file of this patent is not represented in the printed of the file of this patent is not represented in the printed of the file of this patent is not represented in the printed of the printed of this patent is not represented in the printed of the print The invention relates to a novel pharmaceutical composition, which has a at ftp.wipo.int/pub/published_pct_sequences

Sequence 20 BP; 2 A; 6 C; 8 G; 4 T; 0 U; 0 Other;

ö 3.2%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 3.2e+02; Live 0; Mismatches 4; Indel8 125 CGGCATGCTGGCCCGCCTGG 144 1 ceccanerreceasecres 20 16; Conservative Similarity Query Match Local Matches à

Gaps

RESULT 241 AB2987

ABZ98765 standard; DNA; 20 BP

ABZ98765;

17-OCT-2003 (first entry)

Human tryptase b oligonucleotdie sequence.

Human, antisense, lung dysfunction, nasal airway dysfunction, antinflammatory steroid, ubiquinone, antinflammatory, antiallergic, antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy, antisense gene therapy, respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds. 

Homo sapiens.

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Katz E, Pabalan J, Aguilar D; Li Y, Sandrasagra A, K. Tang L, Shahabuddin S; Miller S, Nyce JW,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 14007; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the cinttation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 mucleotides of junctions of genes encoding a polypeptide associated with lung and/or charal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubjourners. A composition of the invention has antiinflammatory, antialergic, antiasthmatic, hypotensive, went an antistlammatory and cytotatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or use in antisense gene therapy. The composition is useful for treating or use in antisense gene therapy. The composition of confiction, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing bronchoconstriction, lung surfactant in a subject's tissue, or treating bronchoconstriction, lung surfactant in a subject's tissue, or treating bronchoconstriction, lung surfactant in a subject's tissue, or arespiratory disease or condition.

Co lung surfactant in a was obtained in electronic format directly from WIPO Ext. The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO Ext. 

Seguence 20 BP; 3 A; 5 C; 7 G; 5 T; 0 U; 0 Other;

Gaps ö 3.2%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 3.2e+02; ative 0; Mismatches 4; Indels 16; Conservative Query Match Best Local Similarity Matches

RESULT 242 ACC62132,

ACC62132 standard; DNA; 20

ACC62132;

20-JUN-2003 (first entry)

Human alipoprotein B antisense oligonucleotide SEQ ID NO: 21

alipoprotein B; ApoB; antilipaemic; antiarteriosclerotic; antidiabetic; anorectic; cardiovascular; gene therapy; lipid metabolism; cholesterol metabolism; atherosclerosis; hyperlipidaemia; diabetes; type 2 diabetes; obesity; atherosclerosis; cardiovascular disease; glucose; antisense oligonucleotide; ss.

Synthetic.

WO2003011887-A2.

13-FEB-2003.

30-JUL-2002; 2002WO-US024247

01-AUG-2001; 2001US-00920033. 30-APR-2002; 2002US-00135985. 15-MAY-2002; 2002US-00147196.

(ISIS-) ISIS PHARM INC.

Crooke RM, Graham MJ;

WPI; 2003-268105/26.

New antisense oligonuclectides for modulating apolipoprotein B, especially for preventing or treating atherosclerosis, hyperlipidemia or diabetes, or for modulating glucose, cholesterol, lipoprotein or trigilyceride levels. 

Example 15; Page 96; 160pp; English

length that is targeted to a nucleic acid molecule encoding to a proper clean that is targeted to a nucleic acid molecule encoding to applipoprotein B (ApoB), and specifically hybridises with and inhibits the expression of a nucleic acid molecule encoding ApoB; or which active site on a nucleic acid molecule encoding ApoB; or which active site on a nucleic acid molecule encoding ApoB. A compound of the anoretic, and cardiovascular activity. The compound may have a use in anoretic, and cardiovascular activity. The compound may have a use in gene therapy. The antisense oligonocleotide is useful for treating an animal having a disease or conditions associated with ApoB. e.g. a condition involving abnormal lipid metabolism, a condition involving abnormal lipid metabolism, atherosclerosis, or a condition involving an abnormal lipid metabolism, atherosclerosis or condition involving an abnormal lipid metabolism, atherosclerosis or a condition involving an abnormal metabolism, aberosclerosis, or a condition involving an abnormal metabolism, absormation (e.g. hyperlipidaemia, diabetes of specifically Type 2 diabetes), obserty, atherosclerosis or cardiovascular disease). The new compound or the antisense oligonucleotide is also useful for modulating glucose levels; (particularly plasma or sernam glucose levels) in a human or diabetic animal, or for modulating serum cholesterol levels, lipoprotein levels (specifically VLDL, HDL or LDL) or serum triglyceride levels, contactive or modulating serum cholesterol levels, lipoprotein levels or preventing or delaying the onset of a disease or condition associated with ApoB, or the onset of an increase in glucose levels in the animal or human. The present sequence is used in the exemplification of the invention relates to a novel compound that is 8-50 nucleotides in

Sequence 20 BP; 5 A; 9 C; 6 G; 0 T; 0 U; 0 Other;

Gaps . 0 Match 3.2%; Score 13.6; DB 1; Length 20; Local Similarity 80.0%; Pred. No. 3.2e+02; es 16; Conservative 0; Mismatches 4; Indels TGCTGGCCCGCCTGGCGGTG 149 130 Query Match Best Loc Matches

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recreececrecreececre 1 20 RESULT 243 ADB25658, 음

fisp-12; NOV2; insulin-like growth factor binding protein-related protein 2; IGFBP-rP2; insulin-like growth factor binding protein-related protein 2; IGFBP-rP2; IGFBP-8; HGSP4; ecogenin; acute lymphoblastic leukaemia; gene therapy; hyperproliferative disorder; cancer; pulmonary fibrosis; renal fibrosis; scleroderma; atherosclerosis; cytostatic; dermatological; antiarteriosclerotic. Human connective tissue growth factor antisense oligo DNA (SeqID 51) antisense; human; ss; connective tissue growth factor; CTGF; chromosome 6g23.1; ctgrofact; fibroblast inducible secreted protein; ADB25658 standard; DNA; 20 BP (first entry) 20-NOV-2003 ADB25658; 

Location/Qualifiers . .20 *tag= nodified base Homo sapiens,

/note= "OTHER= phosphorothioate backbone, where 1-5 and 16-20 are 2' methoxyethyl nucleotides. All cytidines are 5-methylcytidines" 'mod_base= OTHER ದ

WO2003053340-A2.

03-JUL-2003.

09-DEC-2002; 2002WO-US038618.

10-DEC-2001; 2001US-00006191.

(ISIS-) ISIS PHARM INC. 

Watt AT; WPI; 2003-559091/52 Gaarde WA,

New antisense oligonucleotides for modulating connective tissue growth factor expression, particularly useful for treating cancers (e.g. breast or prostate cancer), pulmonary or renal fibrosis, scleroderma or atherosclerosis.

Example 15; Page 85; 139pp; English.

This invention relates to novel methods for modulating the expression of connective tissue growth factor (CTGF) by antisense oligonucleotides.

CTGF has been mapped to human chromosome region 623.1, and is also known as ctgrefact, fibroblast inducible secreted protein, fisp-12, NOV2, insulin-like growth factor binding protein-related protein 2, IGFBF-FP2, IGFBF

Sequence 20 BP; 3 A; 9 C; 4 G; 4 T; 0 U; 0 Other;

ö Score 13.6; DB 1; Length 20; Pred. No. 3.2e+02; 0; Mismatches 4; Indels 80.08; Query Match
Best Local Similarity 80.0
Matches 16; Conservative

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167 GGTGTACTACGAGTCCAAGG 186 20 GGTGTGTGACGAGCCCAAGG 1 ð a

ACD44753 standard, DNA; 20 BP. ACD44753; RESULT 244 ACD44753 

(first entry) 09-SEP-2003 PKA regulatory subunit RII alpha inhibitory oligonucleotide ISIS102782.

Human; se; antisense therapy; infection; inflammation; tumour; protein kinase A regulatory subunit RII alpha.

Homo sapiena, Synthetic

US6524854-B1.

11-SEP-2001; 2001US-00954560. 25-FEB-2003.

11-SEP-2001; 2001US-00954560

(ISIS-) ISIS PHARM INC

Monia BP,

WPI; 2003-511923/48.

rng.res

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inhibitory oligonucleotide
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The invention relates to antisense compounds targeted to nucleic acids encoding protein kinase A regulatory subunit RII alpha. The antisense compounds are useful for modulating the expression of protein kinase A (PKA) regulatory subunit RII alpha and for treating a disease or condition associated with expression of PKA regulatory subunit RII alpha. The compounds are also useful as research reagents and kits, or for diagnostics, therapeutics and prophylaxis, e.g. to prevent or delay infection, inflammation or tumour formation. The present sequence represents a human protein kinase A regulatory subunit RII alpha
New antisense compounds, useful for modulating the expression of protein kinase A (PKA) regulatory subunit RII alpha, and for treating a disease or condition associated with expression of PKA regulatory subunit RII
                                                                                                                                                                                                                                                                                       Claim 14; Col 43-44; 35pp; English
                                                                                                                                                                                   alpha.
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Match 3.2%; Score 13.6; DB 1; Length 20; Local Similarity 80.0%; Pred. No. 3.2e+02; les 16; Conservative 0; Mismatches 4; Indels Sequence 20 BP; 2 A; 8 C; 8 G; 2 T; 0 U; 0 Other; 318 CGCGTGCTGCCGGCGGACGA 337 Query Match Matches

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Gaps

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ADB46018 standard; DNA; 20 Primer #1 of the invention 04-DEC-2003 (first entry) ADB46018; 

protein breakdown; ss; primer. Synthetic.

20-AUG-2002; 2002WO-JP008376 WO2003070954-A1 28-AUG-2003.

21-FEB-2002; 2002JP-00045090

(NODA ) NODA INST SCI RES. (NAAD-) NAT INST ADVANCED IND SCI & TECHNOLOGY.

Umitsuki G, Hatamoto O, Hara S, Masuda T, Sano M, Machida M; WPI; 2003-697623/66 Proteins for increasing breakdown efficiency of protein-containing

Disclosure; Page 61; 77pp; Japanese.

The present invention relates to proteins that have been found useful increasing the breakdown efficiency of protein-containing substances. present sequence represents a primer of the invention.

Sequence 20 BP; 2 A; 7 C; 5 G; 2 T; 0 U; 4 Other;

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Gaps
                                                            ö
Query Match
3.2%; Score 13.6; DB 1; Length 20;
Best Local Similarity 72.2%; Pred. No. 3.2e+02;
Matches 13; Conservative 3; Mismatches 2; Indels
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Identifying biomarker genes using nucleic acid microarrays, useful for molecular diagnostic and pathology applications, comprises comparing the Gibbs-likelihood ratios for each gene and determining a rank order for
                                                                                                                        ss; primer; biomarker gene; gene expression; nucleic acid array; molecular diagnostic method; molecular target.
                                                                                                                                                                                                                                                     Dooley TP, Curto BV, Davis RL;
333
BP.
                                                                                                          COL6Al forward gRT-PCR primer
                                                                                                                                                                                                    10-FEB-2003; 2003WO-US003673.
                                                                                                                                                                                                                     08-FEB-2002; 2002US-0354519P.
                                                         ADC46898 standard; DNA; 20
                                                                                          18-DEC-2003 (first entry)
                                                                                                                                                                                                                                     (INTE-) INTEGRIDERM INC
                                                                                                                                                                                                                                                                     WPI; 2003-731515/69.
                                                                                                                                                                    WO2003067217-A2.
                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                     14-AUG-2003.
                                                                          ADC46898;
                                                                                                                                                                                                                                                                                                               the gene.
                                                   ADC46898
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The invention relates to a method of identifying one or more biomarker genes for a type of cells among a group of (m) different cell types, from a multiplicity of genes whose expression levels in cells of the group are measured using nucleic acid arrays, to generate a plurality of measurements of expression levels for the m types of cells, by comparing rank order for the gene and (m-1) for each gene and detarmining a rank order for the gene among the multiplicity. The method is useful in identifying biomarkers using nucleic acid microarrays. The biomarkers of skin may be used in molecular diagnostic and pathology applications in normal and abnormal tissues and cell. The biomarker genes may also be used as molecular targets for therapeutics of a disorder or a disease in thumans. This sequence represents a QRT-PCR primer used in the method of the invention.

Example 3; Page 38; 54pp; English.

Query Match
3.2%; Score 13.6; DB 1; Length 20;
Best Local Similarity 80.0%; Pred. No. 3.2e+02;
Matches 16; Conservative 0; Mismatches 4; Indels Sequence 20 BP; 5 A; 7 C; 5 G; 3 T; 0 U; 0 Other; 104 TGACCGCGACCGCAGCAAGT 123 1 TGACCCGACCTCAGAGAGT 셤 ò

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RESULT 247
ADE14433/c
ID ADE14433 standard; DNA; 20
XX
AC ADE14433;
XX

WO9618736-A2

20-JUN-1996.

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Arthritic condition; graft tolerance; immune response; target; cleavage; hammerhead xibozyme; hairpin ribozyme; human; rabbit; mouse; collagenase; stromelysin; synovial membrane; joint; arthritis; osteoarthritis; rheumatoid arthritis; autoimmune disease; allergy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention describes a compound (I) 8-80 nucleobases in length targeted to a nucleic acid molecule encoding hydroxysteroid 11-beta dehydrogenase 1. inhibiting expression of hydroxysteroid 11-beta dehydrogenase 1. The methods and compositions of the present invention are useful for treating disorders associated with hydroxysteroid 11-beta dehydrogenase 1 expression, such as osteoporosis, depression and metabolic disorders like obesity, diabetes, atherosclerosis and hyperlipidaemia. This sequence represents an antisense oligonucleotide used to control the expression of human hydroxysteroid 11-beta
                                                                                                                                                                                                                                                                                                                                                                                                                                                  New antisense compounds useful for treating disorders associated with hydroxysteroid 11 beta dehydrogenase 1 expression, such as osteoporosis, depression and metabolic disorders like obesity, diabetes and atherosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Gaps
                                                               osteopathic; antidepressant; anorectic; antidiabetic; antiarteriosclerotic; antilipemic; antisense-therapy; Approxysteroid 11-beta dehydrogenase 1; osteoporosis; depression; metabolic disorder; obesity; HSD11B1; diabetes; atherosclerosis; hyperlipidaemia; antisense technology; human; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .Match 3.2%; Score 13.6; DB 1; Length 20; Local Similarity 80.0%; Pred. No. 3.2e+02; es 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human B7-1 hammerhead ribozyme target SEQ ID NO:1187.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 6 A; 4 C; 6 G; 4 T; 0 U; 0 Other;
                                   HSD11B1 antisense oligonucleotide seq id 35.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 3; SEQ ID NO 35; 53pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CGACTICCICACTITCCIGG 378
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               caactrccrcagrracered 1
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                                                                                                                                                                                                                                                                                                                  19-APR-2002; 2002US-00126355.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
29-JAN-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                   (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-852782/79.
                                                                                                                                                                                                             US2003198965-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      dehydrogenase
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-JUL-1999
                                                                                                                                                                            Homo sapiens.
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                                                                                                                                                                                                                                             23-OCT-2003.
                                                                                                                                                                                                                                                                                                                                                                                        Freier SM;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               359
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The present invention describes a novel enzymatic nucleic acid (ENh) having a hammerhead motif (HM) comprising; (i) at least 5 tibose residues (i) in a 2-C-allyl modification at position 4 of the ENh; (iii) at least ten 2-O-methyl modifications; and (iv) a 3'-end modification. The ENh's can inhibit collagenase and stromelysin production in the synovial membrane of joints for the treatment or prevention of arthritis, particularly osteoarthritis or rheumatoid arthritis. The ENh's can also be used to treat antigen presenting cells of a donor to induce tolerance in a recipient to an alloantigen of a donor. They can also be used for channing graft tolerance or for treating autoimmune disease, and for treating allergies and other inflammatory conditions. The ENh's can also be used in diagnosis Ribozyme therapy impacted on the expression of stromelysin without introducing the non-specific effects upon gene expression which accompany treatment with retinoids and dexamethasone. The present sequence is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Arthritic condition; graft tolerance; immune response; target; cleavage;
                                                                                                                                                                                                                                                                                                                                                                                                                Enzymatic nucleic acid molecules having a hammer-head motif - used for the treatment of arthritis, induction of graft tolerance or treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                   , Pavco P;
Matulic-Adamic J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3.1%; Score 13.4; DB 1; Length 15; 60.0%; Pred. No. 1.9e+02; ative 5; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human B7-1 hammerhead ribozyme target SEQ ID NO:1189.
                                                                                                                                                                                                                                                                                                                   Stinchcomb DT, Jarvis T, Draper K, Gustofson J, Usman N, Wincott F, M Thompson JD, Modak A, Burgin A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 15 BP; 2 A; 3 C; 5 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 10; Page 166; 307pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP
                                                                                       94US-00354920.
94US-00363253.
94US-00363254.
95US-00390850.
95US-00426124.
                                                                                                                                                                     95US-00432874.
95US-00434509.
95US-0000951P.
95US-00512861.
95US-00512861.
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                                                           95WO-US015516
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                                                                                                                                                                                                                                                                                      (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      9; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                  auto-immune diseases.
                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1996-300653/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 9; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       present invention
                                                                                                                                                                                                                                                                                                                     Beigelman L,
Mcswiggen J,
Karpeisky A,
                                                                                                                                                                                                                       07-JUL-1995;
07-AUG-1995;
05-OCT-1995;
                                                                                                                       23-DEC-1994;
17-PEB-1995;
20-APR-1995;
02-MAY-1995;
04-MAY-1995;
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                                                           22-NOV-1995;
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AAX64557
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30-MAR-2001

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hammerhead ribozyme; hairpin ribozyme; human; rabbit; mouse; collagenase; stromelysin; synovial membrane; joint; arthritis; osteoarthritis; rheumatoid arthritis; autoimmune disease; allergy; inflammation; diagnosis; ss.
                                                                                                                                                                                                                                                                                                                        Enzymatic nucleic acid molecules having a hammer-head motif - used for the treatment of arthritis, induction of graft tolerance or treatment
                                                                                                                                                                                                                                                                Stinchcomb DT, Jarvis T, Draper K, Pavco P;
Gustofson J, Usman N, Wincott F, Matulic-Adamic J;
Thompson JD, Modak A, Burgin A;
                                                                                                                                                                                                                                                                                                                                                              Claim 10; Page 166; 307pp; English.
                                                                                                                            94US-00363253.
94US-00363253.
94US-00363254.
95US-00426124.
95US-00432874.
95US-00432874.
95US-000951P.
95US-000951P.
95US-000951P.
                                                                                                             95WO-US015516
                                                                                                                                                                                                                                              (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                  the treatment of arth auto-immune diseases.
                                                                                                                                                                                                                                                                                                      WPI; 1996-300653/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               present invention
                                                                                                                                                                                                                                                                Beigelman L,
Mcswiggen J,
Karpeisky A,
                                                                                                                                                23-DEC-1994;
17-FEB-1995;
20-APR-1995;
                                                                                                                                                                           02-MAY-1995;
04-MAY-1995;
07-JUL-1995;
                                                                       WO9618736-A2
                                                                                                            22-NOV-1995;
                                                                                                                                                                                                        07-JUL-1995
                                                                                                                                                                                                                           05-OCT-1995;
                                                                                         20-JUN-1996
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The present invention describes a novel enzymatic nucleic acid (ENA)
having a hammerhead motif (HM) comprising: (i) at least 5 ribose residues
(ii) a 2-C-allyl modification at position 4 of the ENA; (iii) at least
c in 2'-O-methyl modifications; and (iv) a 3'-end modification. The ENA's
membrane of joints for the treatment or prevention of arthritis,
can inhibit collagenase and stromelysin production in the synovial
membrane of joints for the treatment or prevention of arthritis,
particularly ostecarthritis or rheumatoid arthritis. The ENA's can also
be used to treat antigen presenting cells of a donor to induce tolerance
in a recipient to an alloantigen of a donor. They can also be used for
enhancing graft tolerance or for treating autoimmune disease, and for
treating allergies and other inflammatory conditions. The ENA's can also
be used in diagnosis. Ribozyme therapy impacts of the expression of
stromelysin without introducing the non-specific effects upon gene
expression which accompany treatment with retinoids and dexamethasone.
The concentration of ribozyme required to affect a therapeutic treatment
is lower than that required of antisense molecules, and is highly
concentral to the present sequence is used in the exemplification of the

Seguence 15 BP; 2 A; 4 C; 4 G; 0 T; 5 U; 0 Other;

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Match 3.1%; Score 13.4; DB 1; Length 15; Local Similarity 60.0%; Pred. No. 1.9e+02; es 9; Conservative 5; Mismatches 1; Indels
    Query Match
                          Best Loca
Matches
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402 GICTICTACGIGATC 416 

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AAF53589/c ID AAF53589 standard; DNA; 15 XX RESULT 250

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cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor I receptor; IGF1; pityriaais; IGF binding protein, IGFB-2; IGFBP3; circlammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition; between the condition of the retina; ser
                                                                                                                                                                                                                                                                                                                                                                                                   Ameliorating the effects of a disorder, e.g. psoriasis, by administering W (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation.
                                                                          Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic;
                                                                                                                                                                                                                                                                                                                                                   Edmondson SR;
                                                                                                                                                                                                                                                                                                                         (MURD-) MURDOCH CHILDRENS RES INST.
                                                                                                                                                                                                                                                                                                99US-0140345P.
                                                                                                                                                                                                                                                                       21-JUN-2000; 2000WO-AU000693
                                                 IGF-I oligonucleotide #4549.
                        (first entry)
                                                                                                                                                                                                                                                                                                                                                   Wraight CJ, Werther GA,
                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-041421/05.
                                                                                                                                                                                                                   WO200078341-A1
                                                                                                                                                                                                                                                                                                21-JUL-1999;
                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                              28-DEC-2000.
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The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like drowth Pactor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inhibiting or reducing growth factor mediated cell proliferation, oligonucleotide which can be used to design the antisense oligonucleotide of the present invention (see AAF45151 and AAF45153-61150 une method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, tuba, pilaris, serborrhoea, keloids, keratosis, chothyosis, pityriasis, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic mission of the inside of blood Example 8; Page 90; 201pp; English.

Gaps ö 3.1%; Score 13.4; DB 1; Length 15; 93.3%; Pred. No. 1.9e+02; tive 0; Mismatches 1; Indels Sequence 15 BP; 3 A; 1 C; 7 G; 4 T; 0 U; 0 Other; 14; Conservative Similarity Query Match Best Local 8 Matches

vessels or any other hyperplasia

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AAF84002 standard; DNA; 16 AAF84002; AAF84002

RESULT 251

22-AUG-2001

SAXXX

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The invention relates to a method of treating deficiency of insulin, that involves administering a hedgehog protein or nucleic acid encoding the hedgehog protein. The hedgehog proteins neb eused in the method are selected from sonic hedgehog (Shh), indian hedgehog (Ihh) and desert hedgehog (Dhh). The method is useful for treating deficiency of insulin in a patient afflicted with diabetes, by stimulating insulin production in pancreatic beta-cells PBO). It is also used to treat deficiency of insulin in partient, by stimulating IDX-1 production in PBC. The hedgehog protein is useful for modulating IDX-1 gene expression or its protein in PBC. This is used to treat deficiency of PBC in a patient, by stimulating neogenesis form beta-cell pancreatic ductal precursor cells. Inhibitors of the hedgehog proteins are useful for suppressing secretion of insulin in a patient afflicted with hyperinsulinemia. Sequences AAF84001-4002 represent PCR primers for ampliying the rat Dhh cDNA fragment
                                                                                                                                                                                                                                                                                                                                                                       Treating deficiency of insulin, IDX-1 or pancreatic beta cells in a patient by, administering a hedgehog protein, nucleic acid encoding the protein or cells expressing the protein.
                             Insulin; hedgehog protein; sonic hedgehog; Shh; indian hedgehog; Ihh; desert hedgehog; Dhh; diabetes; panoreatic beta-cell; PBC; IDX-1; neogenesis; hyperinsulinemia; PCR primer; 88.
Rat desert hedgehog (Dhh) cDNA fragment amplifying reverse primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 16 BP; 2 A; 4 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Page 29; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABN07569 standard; DNA; 17 BP.
                                                                                                                                                                                                        08-DEC-2000; 2000WO-US033575.
                                                                                                                                                                                                                                         99US-0170282P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                          (GEHO ) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                             Habener JF, Thomas MK;
                                                                                                                                                                                                                                                                                                                                              WPI; 2001-381492/40.
                                                                                                                                     WO200141786-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-MAY-2002
                                                                                                                                                                                                                                           10-DEC-1999;
                                                                                                                                                                        14-JUN-2001
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                                                                                                        Rattus sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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 BXBXGXGXGXGXGXGXG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ઠે
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Gaps
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% Match 3.1%; Score 13.4; DB 1; Length 16; Local Similarity 93.3%; Pred. No. 2.2e+02; les 14; Conservative 0; Mismatches 1; Indels
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CTGCACTACGAGGGC 80 15 checkchaceaec 1 Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:7561.

Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.

Homo sapiens

WO200192524-A2.

36-DEC-2001

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The present invention describes a human genome-derived myosin-like

protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-

complete acids in gene therapy and vaccine production. The hGDMLP-1

complete acids can be used as probes to detect, characterise and quantify

mucleic acids can be used as probes to detect, characterise and quantify

complete initial substrates for the recombinant engineering of hcDMLP-1

provide initial substrates for the recombinant engineering of hcDMLP-1

complete initial substrates for the recombinant engineering of hcDMLP-1

protein variants having desired phenotypic improvements, and for

expressing the proteins. The hcDMLP-1 proteins or polypeptides may be

completed and a sea standards in assays used to determine the concentration

and/or amount specifically of hGDMLP-1 proteins, as specific biomolecule

complete probes for surface-ahanced laser desorption ionisation, as

therapeutic supplement in patients having specific deficiency in hGDMLP-1

production, and in vaccines or for replacement therapy. The

production, and in vaccines or for replacement therapy. The

production, and in vaccines or for replacement therapy. The

production, and in vaccines or for replacement therapy. The

production, and in vaccines or for replacement therapy. The

complete associated with the expression of hGDMLP-1, in particular heart

and skeletal muscle disorders. hGDMLP-1 may be used for diagnosing a

complete represents an oligomer used in the screening of the

hGDMLP-1 sequence in the exemplification of the present invention. N.B.

The present sequence data for this patent did not form part of the printed

specification, but was obtained in electronic format directly from WIPO

at ftp.wipo.int/pub/published_pct_sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chen W, Shannon ME;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rank DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; SEQ ID NO 7561; 214pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hanzel DK,
                                                                                                                27-SEP-2000; 20000S-0236359P.
04-0CT-2000; 20000S-02036559.
30-JAN-2001; 2001WO-US000661.
30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
                                                                                                                                                                                                                                                                                                                  30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000670.
05-FEB-2001; 2001US-0266860P.
                          25-MAY-2001; 2001WO-US016981
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Su Y, Ji Y, Penn SG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-179446/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (AEOM-) AEOMICA INC.
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3.1%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 2.5e+02; tive 0; Mismatches 1; Indels ... Sequence 17 BP; 6 A; 3 C; 7 G; 1 T; 0 U; 0 Other; 387 GACGCCCCAAGAAG 401 2 gaccececcaacaaca 16 ABN79929 standard; DNA; 17 Query Match
Best Local Similarity 93.3
Matches 14, Conservative RESULT 253 ABN79929/c δ ద

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0; Gaps

15-JUL-2002 (first entry) ABN79929; exxxex

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Query Match 3.1%; Score 13.4; DB 1; Length 17; Best Local Similarity 93.3%; Pred. No. 2.5e+02; Matches 14; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 1 A; 7 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human MDZ3 scanning oligonucleotide SEQ ID 481.
                                                                                                                                                    Ronaghi M, Ekstroem B, Pourmand N;
                                                                                                                    (PYRO-) PYROSEQUENCING AB.
(STRD ) UNIV LELAND STANFORD JUNIOR.
                                                                                                                                                                                                                          Example 2; Page 47; 86pp; English,
                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADA99492 standard; DNA; 17 BP.
                                                                                     10-SEP-2001; 2001WO-GB004042.
                                                                                                      08-SEP-2000; 2000GB-00022069.
                                                                                                                                                                                                                                                                                                                                                                                                       268 ACCTGGAGCAGGGCG 282
                                                                                                                                                                                                                                                                                                                                                                                                                      17 ACCTGGAGCAGAGCG 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-NOV-2003 (first entry)
                                                                                                                                                                    WPI; 2002-393849/42.
                                                                                                                           (STRD ) UNIV LELANI
(GARD/) GARDNER R.
                                                      WO200220837-A2.
                                                                                                                                                                                                           incorporation.
                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EP1281758-A2
                                                                      14-MAR-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADA99492;
                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 25
ADA99492
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The present invention relates to novel human zinc finger-containing corrects and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is proceins and their coding sequences: MDZ3, MDZ4, MDZ12. MDZ3 is encoded at chromosome 7q22.1, MDZ4 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 16p12.2 and MDZ12 is encoded at chromosome 16p2.3 and MDZ3 is encoded at chromosome 16p2.3 and MDZ4, MDZ3, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3. MDZ4, MDZ3, or MDZ12, e.g. cancer or developmental disorders. The nucleic caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids and sloo be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The process are useful as therapeutic agents for gene therapy or as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cytostatic; immunostimulant; gene therapy; vaccine; human;
zinc finger protein; MDZ3; MDZ4; MDZ1; Chromosome 7q22.1;
chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer;
developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 17 BP; 4 A; 3 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human MDZ3 scanning oligonucleotide SEQ ID 479.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 8; SEQ ID NO 481; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADA99490 standard; DNA; 17 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-AUG-2001; 2001US-00922181.
                                                                                                                       30-JUL-2002; 2002EP-00016874
                                                                                                                                                                                                    02-AUG-2001; 2001US-00922181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 292 TGGTGAAGGACCTGA 306
                                                                                                                                                                                                                                                                                                                                                             Shannon M, Gu Y, Nguyen C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          redregaedactrea 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-423107/40.
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                                          05-FEB-2003.
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ADA9949994

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DT 20-NOV

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a novel method for obtaining typing information about several variable sites within target nucleic acid, or typing one or more nucleic acid molecules. The methods of the invention are useful for typing one or more nucleic acid molecules containing two or more variable sites, preferably nucleic acid molecules containing two or more variable sites, preferably nucleic acid molecules containing three or more variable sites are typed, where three or more primer extension reactions are performed. The method is also useful for diagnosis of pathological conditions characterized by the presence of specific nucleic acid microbial species or their subtypes, and in typing procedures e.g. typing of polymorphisms, tissue typing or in clinical applications. The sequence specific target region of genomic DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ1; MDZ1; chromosome 7g22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; se.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Typing nucleic acid for obtaining information about several variable sites involves simultaneously or sequentially performing two or more primer extension reactions, and determining the pattern of nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gapa
                                                                        Human, single nucleotide polymorphism; nucleic acid typing; primer, tissue typing; sequencing; angiotensin converting enzyme; ACE; ss.
Human angiotensin converting enzyme SNP-fragment Eu6 primer A063FS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;
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Gu Y, Nguyen C; WPI; 2003-423107/40. 3P1281758-A2 Shannon M, ADA99413; 256 ADA99413 셤 ò

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The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ12, MDZ12. MDZ3 is encoded at chromosome 7422.1, MDZ4 is encoded at chromosome 6721.3-22.2, MDZ7 is encoded at chromosome 6721.3-22.2, MDZ7 is encoded at chromosome 6721.2 and MDZ12 is encoded at chromosome 15q26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 4 A; 3 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                Example 8; SEQ ID NO 479; 103pp; English.
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Gaps
                                                      .
Query Match 3.1%; Score 13.4; DB 1; Length 17; Best Local Similarity 93.3%; Pred. No. 2.5e+02; Matches 14; Conservative 0; Mismatches 1; Indels
                                                                                                       292 TGGTGAAGGACCTGA 306
                                                                                                                                                        recreaseaccrea 17
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Human MDZ3 scanning oligonucleotide SEQ ID 402. ADA99413 standard; DNA; 17 BP. (first entry) 20-NOV-2003

Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MD23; MD24; MD27; MD212; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.

Homo sapiens

30-JUL-2002; 2002EP-00016874.

02-AUG-2001; 2001US-00922181

(AEOM-) AEOMICA INC.

Shannon M, Gu Y, Nguyen C;

WPI; 2003-423107/40.

New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3,

MDZ4, MDZ7 or MDZ12, e.g. cancer.

Example 8; SEQ ID NO 402; 103pp; English.

The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ12, MDZ3. MDZ3. C encoded at chromosome 7422.1, MDZ4 is encoded at chromosome 6P21.3-22.2, MDZ7 is encoded at chromosome 6P2.3 mDZ7 is encoded at chromosome 15q26.1. The MDZ3, MDZ4, MDZ7, MDZ4 and MDZ12 is encoded at chromosome 15q26.1. The MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or devalopmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as proteins are useful as therapeutic agents for gene therapy or as 

Sequence 17 BP; 2 A; 8 C; 1 G; 6 T; 0 U; 0 Other;

Gaps ö Query Match
Best Local Similarity 93.3%; Pred. No. 2.5e+02;
Matches 14; Conservative 0; Mismatches 1; Indels

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δ

B ADA99491 standard; DNA; 17 ADA99491; RESULT 257 ADA99491

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20-NOV-2003 (first entry)

Human MDZ3 scanning oligonucleotide SEQ ID 480.

Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ12; chromosome 7q22.1; chromosome 6p21;3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss. CCCCX2X444444X4X4X4X6X6X6X6X8X8X8X6X6X6XX6XX6X

Homo sapiens

EP1281758-A2.

05-FEB-2003.

30-JUL-2002; 2002EP-00016874.

02-AUG-2001; 2001US-00922181.

(AEOM-) AEOMICA INC.

Shannon M, Gu Y, Nguyen C;

WPI; 2003-423107/40.

New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MD23, MDZ4, MDZ7 or MDZ12, e.g. cancer.

Example 8; SEQ ID NO 480; 103pp; English.

The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ1, MDZ1, MDZ3, MDZ2, encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome form of the mDZ1 is encoded at chromosome.

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886666666666888
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15q26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The mUZ1s acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
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Sequence 17 BP; 4 A; 3 C; 7 G; 3 T; 0 U; 0 Other;

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                                       Gabs
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/ Match 3.1%; Score 13.4; DB 1; Length 17; Local Similarity 93.3%; Pred. No. 2.5e+02; les 14; Conservative 0; Mismatches 1; Indels
     Query Match
                        Best Loca
Matches
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ADA99412 standard; DNA; 17 BP. ADA99412 

ADA99412;

20-NOV-2003 (first entry)

Human MDZ3 scanning oligonucleotide SEQ ID 401.

Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.

Homo sapiens.

EP1281758-A2.

05-FEB-2003

30-JUL-2002; 2002EP-00016874.

02-AUG-2001; 2001US-00922181.

(AEOM-) AEOMICA INC.

Shannon M, Gu Y, Nguyen C;

WPI; 2003-423107/40.

New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.

Example 8; SEQ ID NO 401; 103pp; English.

The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7922.1, MDZ4 is encoded at chromosome 1592.1, and MDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 15p11.2 and MDZ12 is encoded at chromosome 15p11.2 and MDZ12 is encoded at chromosome 15p12. and MDZ12 is encoded at chromosome 15p12. and MDZ12 is encoded at chromosome 15p12. ADZ2 is encoded at chromosome 15p12. ADZ2 is encoded at chromosome 15p12. The MDZ1, MDZ3, MDZ4, MDZ7, or MZ12, e.g. cancer or developmental disorders. The nucleic acused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12. The probes are

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                                                                                               Gaps
useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                                                                                     Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras; enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV; anti-rheumatic; cancer; AIDS; ss.
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                                                                     Length 17;
                                                                                              1; Indels
                                             Sequence 17 BP; 2 A; 7 C; 2 G; 6 T; 0 U; 0 Other;
                                                                  Query Match
3.1%; Score 13.4; DB 1;
Best Local Similarity 93.3%; Pred. No. 2.5e+02;
Matches 14; Conservative 0; Mismatches 1;
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                                                                                                                                                                                                                                                                                 Human HER2 DNAzyme substrate #597.
                                                                                                                                                                                                        ABZ65140 standard; RNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-MAY-2001; 2001US-0294140P.
06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
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                                                                                                                       363 Trecreating 377
                                                                                                                                     TICCICACIAICCIG 16
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                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                      05-DEC-2002.
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                                                                                                                                                                                   RESULT 259
                                                                                                                                                                                                ABZ65140
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The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-HER2, K-Ras, H-Ras, and HIV acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZS 9889 - ABZ62216, ABZ64544 - ABZ65531, ABZ65220 - ABZ652246, ABZ652246, ABZ65234, ABZ65230 - ABZ652246, ABZ65234, ABZ65331, ABZ65330 - ABZ652246, ABZ65234, ABZ65331, ABZ65330 - ABZ652246, ABZ65331, ABZ65330 - ABZ65324, ABZ65331, ABZ65330 - ABZ652246, ABZ65331, ABZ65331, ABZ65324, ABZ65334, ABZ65331, ABZ65330 - ABZ653246, ABZ65331, ABZ65331, ABZ65330 - ABZ653246, ABZ65331, ABZ65331, ABZ65330 - ABZ6534, ABZ65331, ABZ65331, ABZ6534, ABZ6534, ABZ65331, ABZ65331, ABZ6534, ABZ6534, ABZ65331, ABZ65331, ABZ6534, ABZ6534, ABZ65331, ABZ65331, ABZ6334, ABZ6534, ABZ65331, ABZ65331, ABZ6534, ABZ6534, ABZ65331, ABZ65331, ABZ6534, ABZ6534, ABZ65331, ABZ65331, ABZ6534, ABZ6534, ABZ65331, ABZ65331, ABZ6534, ABZ6534, ABZ65331, ABZ6534, ABZ6534, ABZ65334, ABZ65334, ABZ6534, ABZ6534, ABZ65334, ABZ6534, ABZ6534, ABZ6534, ABZ65334, ABZ65334, ABZ6534, ABZ6544, Match 3.1%; Score 13.4; DB 1; Length 17; Local Similarity 80.0%; Pred. No. 2.5e+02; les 12; Conservative 2; Mismatches 1; Indels Sequence 17 BP; 2 A; 5 C; 6 G; 0 T; 4 U; 0 Other; ribozymes of the invention Query Match Best Loca Matches

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259 CCACGGTGCACCTGG 273

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Example; Fig 8; 123pp; Japanese.
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                                                                                                                                                                                18-JUL-2000; 2000JP-00218039
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AA290684 standard; DNA; 20
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                                                                                                                                                                                                                                      Tsuji M,
                                                                                                                                                                                                                                                                               WPI; 2002-179794/23
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nes 14; Conserv
                                                                                                                                                                                                           (BMLB-) BML INC
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                                                                                             WO200206467-A1
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Iwasaki T;
                                                                                                                          24-JAN-2002
                                                                  Synthetic.
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                                                    Hominidae
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to murine oligonucleotides (ACC62754-ACC68806), which are associated with tumour suppression, tumour reversion, apoptosis and virus resistance. The oligonucleotides are useful as (1) as probes and primers for detecting, identifying, quantifying and/or amplifying nucleic acid, e.g. as one component of gene chip, in vitro as (anti) sense reagents; and (2) for production of recombinant polypeptides. The oligonucleotides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.
                                                                                                                                                                                             Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; murine; tumour suppression; tumour reversion; apoptosis; virus resistance; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease;
                                                                                                                                                                    Murine oligonucleotide associated with tumour supression, SEQ ID 1117.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17 BP; 4 A; 5 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hominidae LDL receptor related DNA sequence #14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 161; 738pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                           Tuijnder M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABL90998 standard; DNA; 19 BP.
                                                                                  BP.
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ACC63870 standard; DNA; 17
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                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                               (MOLE-) MOLECULAR ENGINES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-333167/31.
                                                                                                                                                                                                                                            schizophrenia; ss.
                                                                                                                                                                                                                                                                                                 WO2003025176-A2
                                                                                                                                                                                                                                                                       Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27-MAY-2002
                                                                                                                                         01-JUL-2003
                                                                                                                                                                                                                                                                                                                            27-MAR-2003
                                                                                                               ACC63870;
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ABL90998
ID ABL9
AC ABL9
XX
XX
DT 27-M
XX
DE HOMI
                                                         RESULT 26
ACC63870/
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The present invention describes a method for detecting lipid metabolism errors in patients using as indicators a set of 65 specific low density lipoprotein (LDI) receptor gene mutations. The method can be used in the diagnosis of an inherited predisposition to the development of diseases associated with hyperlipaemia, such as arteriosclerosis and ischaemic heart disease. ABL91141 encodes the LDL receptor given in ABB90525. ABL91142 to ABB9183 represent PCR primers used in the amplification of the receptor gene. ABL909090 to ABL91140 and ABB90445 to ABB90524 represents sequences used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Egashira T, Ishihara
LDL-R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3.1%; Score 13.4; DB 1; Length 19; llarity 82.4%; Pred. No. 3.1e+02; Conservative 1; Mismatches 2; Indels
Hominidae; low density lipoprotein receptor; LDL receptor; detection; lipid metabolic error; hyperlipaemia; mutation; arteriosclerosis; ischaemic heart disease; ischaemia; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 19 BP; 5 A; 4 C; 7 G; 2 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human KVLQT1 exon 5/intron 5 junction sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Okada T, Nagano M,
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Splawski

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The invention relates to KVLQT1 and KCNEI genes, associated with long QT (LQT) syndrome. It provides a mink protein comprising a mutation which substitutes the wild type amino acids with Leu, Asp, Leu, His, Trp and Ala or Thr at residues 74,76,28,32,98 and 127 respectively. Screening KVLQT1 and KCNEI is useful for identifying mutations for diagnosing and treating LQT. The ability to predict LQT enables physicians to prevent the diseases with medical therapy such as beta blocking agents and opts for better treatments. Sequences AAZ90675-Z90706 represent human KVLQT1
                                                                                                                                            encoding minK protein and KVLQT1 protein involved channel formation useful for screening drugs, for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 2 A; 8 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                    in cardiac potassium channel formation use:
preventing and treating cardiac arrhythmia
                                                                                                                                                                                                                                                                                                     Example 11; Page 69; 167pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       intron/exon junction sequences
Sanguinetti MC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 93.35
                                                                                                                                                forms of genes
                                                                        WPI; 2000-195262/17.
    Keating MT,
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3.1%; Score 13.4; DB 1; Length 20; 93.3%; Pred. No. 3.5e+02; ive 0; Mismatches 1; Indels
                                                                136 CCCGCCTGGCGGTGG 150
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AAZ98914 standard; DNA; 20 06-JUN-2000 AAZ98914; RESULT 263 AAZ98914/c

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Human long QT syndrome-associated KVLQT1 exon 5/intron 5 boundary.
       (first entry)
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potassium channel; KCNEl; ss; Long QT syndrome; gene therapy; KVLQT1; mutation; human; cardiac I(k8)
cardiac arrhythmia; electrocardiogram;
chromosome llpl5.5; intron; exon. location/Qualiflers ď 1. .10 sapiens Homo exon

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/number= 5
11. .20
/*tag= b
                            'number=
           intron
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99WO-US010260 WO200006199-A1 12-MAY-1999; 10-FEB-2000

98US-0094477P (UTAH ) UNIV UTAH RES FOUND. (GENZ ) GENZYME CORP. 29-JUL-1998; 17-AUG-1998;

Landes GM, Curran ME, Sanguinetti MC, Keating MT, Sanguine Burn TC, Splawski I;

Connors TD;

WPI; 2000-195199/17.

New isolated mutant KVLQT1 nucleic acids, useful for developing products for the diagnosis, prevention and treatment of long QT syndrome.

Example 11; Page 72; 178pp; English

The invention relates to KVLQT1 nucleic acids which have a mutation compared to wild-type KVLQT1 (AAZ98901) The KVLQT1 gene encodes a protein of 676 amino acids which forms a cardiac I(ks) potassium channel with the KCNE1 protein (AAY80863). The KVLQT1 gene contains 15 introns and encodes a protein containing 6 putative transmembrane segments and a pore forming region. The gene has been mapped to the chromosomal location 11p15.5. The sequences AAZ8806729836 represent the intron-exon boundaries from the KLVLQT1 genomic sequence. Mutations in the KVLQT1 or KCNE1 genes result in cardiac arrhythmias observed as a prolonged QT curve in electrocardiograms (Long QT syndrome). The genes and proteins can be used to streen for drugs which can be used for treating or preventing long QT syndrome. They can also be used to screen for drugs which can be used for treating or preventing long QT syndrome. The KVLQT1 nucleic acids can be used for gene therapy, and KVLQTI peptides can be used for peptide therapy

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Sequence 20 BP; 2 A; 8 C; 8 G; 2 T; 0 U; 0 Other;
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Gaps . 3.1%; Score 13.4; DB 1; Length 20; 93.3%; Pred. No. 3.5e+02; ive 0; Mismatches 1; Indels Conservative Local Similarity 14; Query Match Matches

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136 CCCGCCTGGCGTGG 150
                     15 CCCACCTGGCGGTGG
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Gaps

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Human PARP-3 antisense inhibitor ISIS #126076. BP. AAS45876 standard; DNA; 20 (first entry) 18-DEC-2001 AAS45876; AAS45876

Human; 88; PARP; Poly (ADP-ribose) polymerase; antisense oligonucleotide; cytostatic; nootropic; neuroprotective; antiinflammatory; antidiabetic; immunosuppressant; hyperproliferative disorder; cancer; cellular injury; oxidative stress; neurological disorder; parkinsonism; apoptosis; meningitis-associated intracranial complication; ischaemia; probe; inflammatory disorder; autoimmune disorder; arthritis; diabetes.

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'mod_base= OTHER
'note= "All cytidine residues are 5-methyl cytidine"
                                                                                                                                       'note= "2'-methoxyethyl nucleotides'
                                                                                           /mod_base= OTHER
/note== "Phosphorothicate backbone"
                                                                             Location/Qualifiers
                                                                                                                                 base= OTHER
                                                                                                    ...20
*tag= b
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                                                                                                                                  pou/
                                                                                  modified base
                                                                                                     modified_base
                                                                                                                         modified base
                                                                    Homo sapiens
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WO200164955-A1.

/*tag= d /mod_base= OTHER /not== "2' methoxyethyl nucleotides"

16. .20

modified_base

07-SEP-2001,

01-MAR-2001; 2001WO-US006572.

Wed Apr 21 12:58:21 2004

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The invention relates to antisense oligonuclectides targeted to human PARP nucleic acid and inhibiting expression of human PARP. PARP (Poly And Andrews) polymerases plays an important role in chromatin decondensation, DNA replication, DNA repair, gene expression, malignant transformation, cellular differentiation and apoptosis. The antisense oligonuclectide inhibitors are useful for inhibiting the expression of PARP in human cells or tissues. They are also useful for treating a human with a disease associated with PARP especially hyperproliferative disorders (e.g. cancer), cellular injury resulting from oxidative stress, environgacial (e.g parkinsonism, meningitis-associated intracranial complications and ischaemia), inflammatory and autoimmune disorders (e.g arthritis) and diabetes. The present sequence is an antisense oligonuclectide of the invention
                                                                                                                                                                                                                                                                                                                                                                                                       Antisense compound useful for treating hyperproliferative, neurological, inflammatory and autoimmune disorders and diabetes inhibits human PARP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 3 A; 7 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 3; Page 91; 168pp; English
02-MAR-2000; 2000US-00517467.
                                                                                                            (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                              Popoff I, Cowsert LM;
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3.1%; Score 13.4; DB 1; Length 20; 93.3%; Pred. No. 3.5e+02; tive 0; Mismatches 1; Indels
                                                                  273 GAGCAGGCGCCACC 287
                                                                                            SAGCAGGGCTGCACC 15
                                  14; Conservative
                Best Local Similarity
Matches 14; Conserv
      Query Match
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Human KVLQTI exon/intron boundary for exon #5
     AAC89924 standard; DNA; 20 BP
                   (first entry)
                   08-MAR-2001
            AAC89924;
   AAC8992
RESULT
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Human, KVLQT1; antiarrhythmic; cardiant; gene therapy; cardiac potassium channel; Jervell and Lange-Nielsen Syndrome; JLN; chromosome 11p15.5; long QT syndrome; ss.

Homo sapiens JS6150104-A.

21-NOV-2000

97US-00874655. 13-JUN-1997; 29-JUL-1998;

98US-00135021

17-AUG-1998;

(UTAH ) UNIV UTAH RES FOUND.

Splawski I; Keating MT,

WPI; 2001-060013/07.

DNA encoding for a mutant KVLQT1 which causes Jervell and Lange-Nielsen syndrome (JLN) when homozygous, useful for diagnosing long QT syndrome, or diagnosing or prognosing JLN.

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Example 5; Col 45-46; 58pp; English.
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KVLQT1 is a cardiac potassium channel and mutations in the KVLQT1 gene cause Jervell and Large-Nielsen Syndrome (JLN). KVLQT1 maps to chromosome 11p15.5. The present invention relates to a mutant KVLQT1 coding sequence (see AAC89914). The mutant KVLQT1 coding sequence is useful in the presence diagnosis of long OT syndrome and in screening humans for the presence of KVLQT1 gene variants which cause JLN syndrome. The present sequence is an exon/intron boundary of KVLQT1 XXX0000000XXX

Sequence 20 BP; 2 A; 8 C; 8 G; 2 T; 0 U; 0 Other;

ö Gaps ö Length 20; 1; Indels Match 3.1%; Score 13.4; DB 1; Local Similarity 93.3%; Pred. No. 3.5e+02; nes 14; Conservative 0; Mismatches 1; Query Match

ò g RESULT 266

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ВЪ.
     AAI69777 standard; DNA; 20
                                                                13-DEC-2001 (first entry)
                                      AA169777;
AA16977
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Bacterium detection; 168/23SrRNA spacer region; PCR primer; 88. 16S/23SrRNA spacer region PCR primer #3.

Pseudomonas putida JP2001190279-A.

ö

Gaps

17-JUL-2001. 

13-JAN-2000; 2000JP-00004160.

(MITO ) MITSUBISHI JUKOGYO KK 13-JAN-2000; 2000JP-00004160.

Detection method of Pseudomonas bacteria. WPI; 2001-605311/69.

المناب والإنام والمارية

Claim 9; Page 8; 11pp; Japanese.

The present invention relates to a method for the detection of the 163/23SrRNA spacer region of Pseudomonas putida (see AAI69774). The method can be used to detect Pseudomonas bacteria. The present sequence is a PCR primer which was used in an example from the present invention

Sequence 20 BP; 4 A; 6 C; 4 G; 6 T; 0 U; 0 Other;

ö 3.1%; Score 13.4; DB 1; Length 20; 93.3%; Pred. No. 3.5e+02; ive 0; Mismatches 1; Indels Best Local Similarity 93.3%; Matches 14; Conservative Query Match

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Gaps

3 CCAGGAGTGAACTG 17 ហ 13

à

AAL40401 standard; DNA; 20 BP. RESULT 267 AAL40401 ID AAL4 XX AAL4

AAL40401

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The invention relates to an antisense oligonucleotide compound of 8 to 50 nucleotides in length that is targeted to a nucleic acid molecule encoding caspase 6, where the oligonucleotide specifically hybridises with and inhibits the expression of caspase 6. The oligonucleotide of the invention specifically hybridises to and inhibits expression of caspase 6 invention specifically hybridises to and inhibits expression of caspase 6 the rerapeutically or prophylactically to treat an animal having a disease or condition associated with caspase 6, such as Rieger's syndrome or axaxia telangiecrasia, hyperproliferative disorder, a haematopoietic disorder, a bone metabolism or cholesterol disorder, various types of cancer, neurological conditions such as Alzheimer's disease and other deregulated apoptoric pathological conditions. This polynucleotide sequence represents a mouse caspase 6 oligonucleotide sequence has 2' MOE wings and NOTE: This phosphorothioate oligonucleotide sequence has 2' MOE wings and
                                                                                                                                                             Muscular; cytostatic; nootropic; neuroprotective; ophthalmological; antilipaemic; osteopathic; caspase 6; Rieger's syndrome; bone metabolism; ataxia telangiectasia; hyperproliferative disorder; cholesterol disorder; haematopoletic disorder; cancer; neurological; Alzheimer's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           An antisense oligonuclectide of 8 to 50 nucleotides in length that inhibits caspase 6, is useful for treating Rieger's syndrome.
                                                                            caspase 6 antisense inhibition related oligo SEQ ID No 120.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 3 A; 5 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Watt AT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 3; Page 92; 141pp; English
                                                                                                                                                                                                                                                                                                                                                           apoptotic; mouse; murine; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-OCT-2001; 2001WO-US030871
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-OCT-2000; 2000US-00679299.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Brown-Driver VL, Zhang H,
19-SEP-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-471315/50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200229066-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     a deoxy gap
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-APR-2002
                                                                                          Mouse
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3.1%; Score 13.4; DB 1; Length 20; 93.3%; Pred. No. 3.5e+02; tive 0; Mismatches 1; Indels
                               14; Conservative
                  Similarity
     Query Match
                  Local
                              Matches
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122 GTACGGCATGCTGGC 136
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ABI94283 standard; DNA; 20 BP 16-FEB-2002 ABI94283; RESULT 268 ABI94283/c 

Capture oligonucleptide Zip ID#1370 oligo #9. (first entry)

Human; K-ras; PCR primer; probe; capture probe; mutation detection; ligase detection reaction; LDR; p53; BRCAL; BRCA2; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndrome; obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forensic;

environmental monitoring; food industry; feed industry;

Synthetic 

WO200179548-A2

25-OCT-2001

04-APR-2001; 2001WO-US010958

14-APR-2000; 2000US-0197271P.

(CORR ) CORNELL RES FOUND INC.

Favis R, Kliman R; Gerry NP, Barany F, Zirvi M,

WPI; 2002-034366/04

Designing capture oligonucleotide probes for use on a support to which complementary oligonucleotides hybridize with little mismatch.

Example 5; Fig 29; 300pp; English.

The present invention describes a method (M1) for designing capture oligonuclectide probes (I) for use on a support to which complementary oligonuclectide probes (II) will hybridse with little mismatch, where coligonuclectide probes (II) will hybridse with little mismatch, where coligonuclectide probes (II) will hybridse with little mismatch, where to algonically little mismatch, where coligonuclectide is useful for detecting infectious diseases caused by bacterial infectious agents confidencially little fundatus, viruses e.g. T-cell lymphocytotrophis cirus, peptential ratus, virus and polio virus, and parasitic infectious agents confidencial many virus and polio virus, and parasitic infectious agents confidencial ratus wolvulus, bitaneoba histolytica and Dracunculus medinesis. The method is also useful for detecting genes, or genes concert in specifically associated with a gene selected from BRCA1 genes involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCA1 gene, muchois also used for environmental monitoring, forensics and the food and feed industry, detecting comprises scanning (using e.g. a scanning electron microscope and infrared microscope) the support at the control of particular sites and infrared microscope) the support at the presence or absence of the target uncleotide sequences. ABI82074 to represent or include sequences used in the exemplification. the present invention

Sequence 20 BP; 2 A; 6 C; 6 G; 6 T; 0 U; 0 Other;

Gaps ö Query Match
3.1%; Score 13.4; DB 1; Length 20;
Best Local Similarity 93.3%; Pred. No. 3.5e+02;
Matches 14; Conservative 0; Mismatches 1; Indels

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175 ACGAGTCCAAGGCAC 189 16 Accadrccaagccac 2 셤

ABZ91337 standard; DNA; 20 RESULT 269 ABZ91337/c 

BP.

17-OCT-2003 (first entry)

Human oligonucleotide sequence.

Human, antisense; lung dysfunction, nasal airway dysfunction, antiinflammatory; antiallergic; antianflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity;

(INRM ) INSERM INST NAT SANTE & RECH MEDICALE.

22-MAR-2001; 2001WO-IB000546. 22-MAR-2001; 2001WO-IB000546

WO200277228-A1

3-OCT-2002

Homo sapiens.

Fischer A;

De Villartay J, Moshous D,

WPI; 2003-029937/02.

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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a cuse in atrisense gene therapy. The composition may have a cuse in atrisense gene therapy. The composition may have a cuse in atrisense gene therapy. The composition may have a cuse in atrisense gene therapy. The composition may have a cuse in a missense gene therapy. The composition may have a cuse in a prophylactic or therapeutic respiratory effect of an antifinanmatory steroid in a subject, for reducing or depleting levels of a ceptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine creceptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject is tissue, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition. Once: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO cases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                                                                    Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; Artemis gene; DNA repair factor; metallo beta-lactamase; RS-SCID; chromosome 10; severe combined immunodeficiency; SCID1; cancer; PCR;
adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                   Li Y, Sandrasagra A, Katz B, Pabalan J, Aguilar D;
Tang L, Shahabuddin S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3.1%; Score 13.4; DB 1; Length 20; 33.3%; Pred. No. 3.5e+02; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer used to amplify Human Artemis gene exon 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 3 A; 10 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; SEQ ID NO 6579; 872pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        l Similarity 93.3%; Pr
14; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP
                                                                                                                                                                                            23-APR-2002; 2002WO-US013135.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           282 GGCACCAAGCTGGTG 296
                                                                                                                                                                                                                                        24-APR-2001; 2001US-0286137P.
                                                                                                                                                                                                                                                                             EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABV72389 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GGCACCAGGCTGGTG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-229219/22
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                                                                                                                WO200285308-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-JAN-2003
                                                                        Homo sapiens.
                                                                                                                                                        31-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ubiquinone
                                                                                                                                                                                                                                                                                                                                           Miller S,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                       Nyce JW,
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Matches
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                                                                                                                                                                                                                                                                           PCR primers ABV72389-ABV72416 were used to amplify exons of the human Artemis gene. This gene encodes a V(D)J recombination and/or DNA repair factor that belongs to the metallo beta-Lactamase superfamily, and whose mutations give rise to the human RS-SCID condition. The gene is localised to chromosome 10. The Artemis gene or its nucleic acid is useful for diagnosing or treating severe combined immunodeficiencies (SCIDS) or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CNS; conserved non-coding region; 88; cytokine; interleukin 4; IL-4; interleukin 5; IL-5; interleukin 13; IL-13; chromosome 5g31; LCR; PCR; locus control region; interleukin gene cluster; transcription factor; human; mouse; dog; rat; bovine; pig; rabbit; fruitfly; puffer fish;
                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                           New isolated nucleic acid molecule of the Artemis gene, useful for diagnosing or treating SCID or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
3.1%; Score 13.4; DB 1; Length 20;
Best Local Similarity 93.3%; Pred. No. 3.5e+02;
Matches 14; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 2 A; 7 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                    Example 1; Page 62; 71pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABX75395 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-FEB-2001; 2001US-00789529
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18-FEB-2000; 2000US-0183657P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Forward PCR primer for CNS-6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             50 CCACTCAGAGGAGTC 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20 CCAATCAGAGGAGTC 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-MAR-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Drosophila melanogaster
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rattus norvegicus.
Oryctolagus cuniculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          primer, transgenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mus musculus.
Canis familiaris.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          JS2002132290-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sus scrofa.
Bos taurus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fugu ripes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABX75395;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 271
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The invention relates to an isolated mucleic acid molecule comprising a length of about 100 nucleotides of less, which has a sequence at least about 100 nucleotides of less, which has a sequence at least about 100 nucleotides of less, which has a sequence (CNS) 1 sequence (a locus control region (CCS) element in interleukin gene cluster region of chromosome 5431 containing interleukin (IL) 4, IL5 and IL 131 . Optionally, the nucleic acid has 70% identity to a human CNS-2 to CNS-16 or mouse CNS-16 or their complements. Also included are: (1) an expression cassette comprising a CNS-1 sequence operably linked to a promoter which controls transcription of a heterologous coding sequence. (2) an expression cassette consisting a Sesentially of an IL-4 gene, an IL-13 gene, and A CNS-1 sequence flanked between two comprising one of the expression cassettes or the T-cell; and (7) a comprising one of the expression cassettes or the T-cell; and (7) a comprising one of the expression cassettes or the T-cell; and (7) a non-human transgenic animal where a CNS-1 sequence, (8) a non-human transgenic compact and lacking a CNS-1 sequence is deleted from its chromosome. The T cell is useful for identifying a compound that computes on mutable procession cassettes or the T-cell; and (7) a non-human transgenic animal where a CNS-1 sequence of the CNS-1 sequence of the compound on binding of the transcription factor to a contacting the compound on binding of the transcription factor or the CNS-1 sequence. The compound on binding of the transcription factor or compound. The nucleic acid is useful for modulate sequence of the CNS-1 sequence or or stroking gene expression. The expression cassettes is custofied to cytoking gene expression. The expression cassettes is a cytoking gene expression. The expression cassettes is a cytoking gene expression. Expression assettes is a in vivo compound. The rushing two lines of non-human transgenic animals that the transcription are intentical except one line has the CNS-1 sequence and the other line
                                                                                                                                                                                                                                Novel isolated nucleic acids which are locus control region elements in interleukin gene cluster region of chromosome, referred as conserved non-coding sequences, useful for modulating expression of cytokine genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       degenerate PCR primer used to isolate a CNS sequence from a variety of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               various therapeutic modalities. The present sequence is a
                                                                                                                                                                                                                                                                                                                                              Example 1; Page 20; 48pp; English.
                                                                                                                                    Loots GG;
                                                                                                                              Frazer KA, Rubin EM,
                                                                                                                                                                                     WPI; 2003-165733/16.
                    (FRAZ/) FRAZER K A.
(RUBI/) RUBIN E M.
(LOOT/) LOOTS G G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               models for
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0; Gaps 3.1%; Score 13.4; DB 1; Length 20; 82.4%; Pred. No. 3.5e+02; ive 0; Mismatches 3; Indels Sequence 20 BP; 5 A; 1 C; 7 G; 5 T; 0 U; 2 Other; 82.48; Conservative Local Similarity 14; Query Match Matches

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HLA; polymerase chain reaction; inflammatory arthropathy; susceptibility; arthritis; arthritis related diseases; ss.

Human leukocyte antigen probe.

WO9207956-A1

Synthetic.

(revised)
(first entry)

25-MAR-2003 19-NOV-1992

AAQ24900;

AAQ24900 standard; DNA; 18 BP.

RESULT 273 AAQ24900

'533/c AAD47533 standard; DNA; 20 BP. RESULT 272 AAD47533/c 2XXXXXXXXX

24-FEB-2003 (first entry)

Human Artemis exon 1 amplifying PCR primer, Ex1F1.

(BRBI-) BRITISH BIO-TECHNOLOGY LTD.

WPI; 1992-183691/22

Hill AV;

90GB-00024005. 91WO-GB001935.

05-NOV-1990;

05-NOV-1991;

14-MAY-1992.

Human, ARTEMIS protein, V(D)J recombination, DNA repair; gene therapy, severe combined immunodeficiency; SCID; cancer; exon 1; PCR; primer; ss.

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The invention relates to an Artemis nucleic acid coding for a protein involved in V(D)J recombination and/or DNA repair. Sequences of the invention are useful for treating severe combined immunodeficiencies (SCID) or cancer. They are also useful for diagnoshing a patient, including a prenatal diagnoshs with SCID, a predisposition to cancer, immune deficiency or a carriage of a mutation increasing the risk of progeny to have such a disease. Peptides of the invention are used for preparing antibodies. The invention is useful in gene therapy. The present sequence is a PCR primer used to amplify human Artemis exon 1 DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Gaps
                                                                                                                                                                                                                                                                                                          New ARTEMIS nucleic acid coding for a protein involved in V(D)J recombination and/or DNA repair, useful for treating and diagnosing severe combined immunodeficiencies (SCID) or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3.1%; Score 13.4; DB 1; Length 20; 93.3%; Pred. No. 3.5e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 2 A; 7 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                    (INRM ) INSERM INST NAT SANTE & RECH MEDICALE.
                                                                                                                                                                                                                                        De Villartay J, Moshous D, Fischer A;
                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 66; 71pp; English.
                                                                                                                                                                22-MAR-2001; 2001WO-IB000546.
                                                                                                                            21-MAR-2002; 2002WO-IB001737
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            50 CCACTCAGAGGAGTC 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20 CCAATCAGAGGAGTC 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity 93.3
Matches 14; Conservative
                                                                                                                                                                                                                                                                            WPI; 2003-018886/01.
                                                    WO200277026-A2.
                  Homo sapiens.
                                                                                      03-OCT-2002.
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25-MAR-2003
05-OCT-1994
                                                  Synthetic.
                                      AAQ56855;
                                                                      viruses.
                                   AAQ56855,
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Sequence 18 BP; 2 A; 7 C; 5 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          recognition site; ds
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14-DEC-1990;
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06-NOV-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
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                                                                                                                                                                                                                                                                                                                  AAQ87132;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAQ92473/C
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                                                                                                                                                                                                                                        RESULT
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                                                                                                                          The sequence is that of a probe which hybridises to one of the human leukocyte antigen (HLA) sequences in the primer extension products (or strands) produced during PCR amplification of the HLA class I alleles. It is specific for the sequence encoding amino acids 67-71 (CXAKA) of the alpha 1 domain of the HLA-B27 group and is thus specific only for this group. It can be used in the detection and/or identification of an HLA sequence that may be indicative of a patients succeptibility to inflammatory arthropathy such as arthritis and arthritis related diseases. Such diseases include reactive arthritis, rheumatoid arthritis, Reiter's syndrom, uveitis, viral arthritis, psoriatic arthropathy, gouty arthritis, septic arthritis, erythema nodosum, Henoch-Schloelein purpura anthritis, septic arthritis. See also AAQ24895-Q24902. (Updated on 25-WAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sets of PCR primers (see AAQ56835-Q56857) are used as probes to detect Norwalk-related viruses, e.g. SRSV/KY/89, HuCV Sapporo, HuCV Houston and primate calcivirus. Detection of viral RNA is by RT-PCR. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA from Norwalk and related viruses - used for preparing prods. for use in diagnostic assays, detection and vaccines for Norwalk and related
                amplification of nucleic acids using buffer soln. and chelating agent or detecting HLA class I alleles for determining susceptibility to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Norwalk virus; HuCV; Sapporo; pathogen; acute gastroenteritis; food poisoning; seafood contamination; diagnostic assay; PCR primer; human calcivirus; small round virus; polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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3.1%; Score 13.2; DB 1; Length 18;
Best Local Similarity 83.3%; Pred. No. 3e+02;
Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primer P-74 for detection of Norwalk-related virus.
                                                                                                                                                                                                                                                                                                                                                                                                                   Seguence 18 BP; 6 A; 6 C; 5 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Graham DY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 49; Page 104; 156pp; English.
                                                                                                   Disclosure; Page 13; 52pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Jiang X,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (BAYU ) BAYLOR COLLEGE MEDICINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       173 CTACGAGTCCAAGGCACA 190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ56855 standard; DNA; 18 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CTGCAAGGCCAAGGCACA
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(first entry)
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arthritis etc.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cleavage of resistant DNA sites with restriction enzymes - using activator comprising recognition site and cleavage-permitting flanking
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                             Gaps
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Score 13.2; DB 1; Length 18;
Pred. No. 3e+02;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA cleavage; restriction endonuclease; NaeI; activator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 18 BP; 0 A; 5 C; 10 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure, Col 21; 23pp; English.
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                                                                                                                       271 TGGAGCAGGGCGCCACCA 288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NaeI substrate oligonucleotide
                                                                                                                                                                                                                                                                                                                  ВЪ.
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        Query Match
Best Local Similarity 83.3%;
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                     AAQ87132 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                        (revised)
(first entry)
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Best Local Similarity 83.3
Matches 15; Conservative
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(first entry)

93JP-00260984 93JP-00260984

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Distinguishing different human herpes virus strains
                                                                                                                                                                                                                       Seguence 18 BP; 4 A; 6 C; 4 G; 4 T; 0 U; 0 Other;
                   Cytomegalovirus detection oligonucleotide #3.
                                                                                                                                                                                                                                                                                                                                                    Human herpesvirus group B primer #1
                                                                                                                                                                   Claim 1; Page 9; 10pp; Japanese.
                                                                                                                                                                                                                                                              216 AACTCGGTGGCGGCCAAA 233
                                                                                                                                                                                                                                                                                                                                                                              sandwich hybridisation; ss
                                                                                                                                                                                                                                                                                                              AAT01523 standard; DNA; 18
                                                                                                                                                                                                                                     Query Match
Best Local Similarity 83.3
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1995-370480/48
                                                                                                                                  WPI; 1995-196320/26.
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                                                                                                                     (TOYM ) TOYOBO KK.
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                                                                                          19-OCT-1993;
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                                                                JP07111893-A.
                                                                                                       19-0CT-1993;
     12-JAN-1996
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                                                                             02-MAY-1995.
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                                                    Synthetic.
                                                                                                                                                                                                                                                                            18
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(first entry)

94JP-00041101 94JP-00041101

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Primers and probes AAT01515-40 and AAT16978-87 are used in a novel method for the specific detection of human herpes viruses (HHV) in which at pleast two types of HHV nucleic acids are pre-amplified by at least 4 primers, followed by a separate detection step using specific detection probes. The primers and probes are synthesised based on the sequences of at least 8 HHV strains selected from HSV1, HSV2, VZV, EBV, CMV, HHV-6A, HHV-6B and HHV-7. They are split into 3 groups: A, B or C. Similarly the probes are split into 3 groups: A, B' and C'. The probes are specific in that they will only detect the amplification prods. from that virus by sandwich hybridisation. This primer is derived from Epstein-Barr virus by and cytomegalovirus (CMV) sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           An oligonucleotide for the amplification and the specific detection of Epstein-Barr virus (EBV) and cytomegalovirus (CMV) - useful for detection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Q876296-Q876303 are PCR primers used in a new method for the amplification and specific detection of Epstein-Barr virus (EBV) and cytomegalovirus (CMV). The oligonucleotides are useful for the detcting the EBV and CMV genes from a culture supernatant of herpes virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
amplification with at least 4 primers and hybridisation to specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Primer; oligonuclectide; Epstein-Barr virus; cytomegalovirus; CMV; amplification; detection; herpes; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Epstein-Barr virus (EBV) and cytomegalovirus (CMV) PCR primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 3.1%; Score 13.2; DB 1; Length 18; Best Local Similarity 83.3%; Pred. No. 3e+02; Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18 BP; 4 A; 6 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                               Sequence 18 BP; 4 A; 6 C; 4 G; 4 T; 0 U; 0 Other;
                                                                     Claim 3; Page 10; 14pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 216 AACTCGGTGGCGGCCAAA 233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  216 AACTCGGTGGCGGCCAAA 233
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ87296;
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ID AAO8
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                                                                                                                  Cytomegalovirus; hybridisation assay; radioisotope; fluorescent compound; enzyme; linker arm; biotin; RNA polymerase promoter; immobilisation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The oligonucleotides AAQ92471-86 can be used for the detection of cytomegaloviruses in a hybridisation assay. The oligonucleotides may be modified by labelling with radioisotopes, fluorescent compounds, enzymes, nucleotides with linker arms, biotin or the promoter sequence for an RNA polymerase. The oligonucleotides may be optionally immobilised
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Primer, PCR; amplification, probe; human, herpes virus; cytomegalovirus; herpes simplex virus; varicella zoster virus; Epstein-Barr virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 modified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Oligo:nucleotide(s) for detection of cytomegalovirus - can be with labels, useful in hybridisation assays, opt. immobilised.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3.1%; Score 13.2; DB 1; Length 18; 83.3%; Pred. No. 3e+02; 1ve 0; Mismatches 3; Indels
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96US-00611280
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                                          Grossman A,
                     (AMGE-) AMGEN CANADA INC
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                                                               WPI; 1996-477128/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    05-NOV-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-NOV-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9820165-A2
                                                                                                                                                                                                                                                                                                                                                                                                                         24-MAR-1999
 03-APR-1996;
                                          Matsuyama T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-MAY-1998.
                                                                                                                                                                                                                                                                                                                                                                                                  AAX10087;
                                                                                                                                                                                                                                                                                                                                                        RESULT 281
                                                                                                                                                                                                                                                                                                                                                                 AAX10087
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                                                                                                                                                                                                                                                                                                                      셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                        AAQBB011-30 are oligonuclectides used for the detection of Epstein-Barr virus. There is no cross reaction with other type of herpes viruses using these probes
                                                                                                                                                                                                                                                                                                                                   Oligo:nucleotide(s) for detection of Epstein-Barr virus - have no cross reactivity with other herpes viruses.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lymphocyte specific interferon regulatory factor; LSIRF; IRF-3; probe; major histocompatibility complex; MHC; ISRE; interferon-stimulated response element; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;
                                                                                                                                 Oligonucleotide probe 10 for detection of Epstein-Barr virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.1%; Score 13.2; DB 1; Length 18; larity 83.3%; Pred. No. 3e+02; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 18 BP; 4 A; 6 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mouse MHC ISRE binding sequence mutant mt4.
                                                                                                                                                         probe; detection; Epstein-Barr virus; ss.
                                                                                                                                                                                                                                                                                                                                                                      Claim 1; Page 9; 10pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      216 AACTCGGTGGCGGCCAAA 233
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                                                                  AAQ88020 standard; DNA; 18 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18 ACCITGGTGGTGGCCAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAT41713 standard; cDNA; 18
 18 ACCTTGGTGGTGGCCAAA
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                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                               WPI; 1995-157847/21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                         (TOYM ) TOYOBO KK.
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                                                                                                                                                                              Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                         AAQ88020;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 280
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Matches
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AAX09121-X10268 are allele-specific oligonucleotide primers used in the isolation of various biallelic polymorphic markers found in the human genome (represented in AAX10269-X12937). These primers can be used in a
                                                                                                                                                                                                                                                                                                                                                                        forms (AAT41710-13) of the murine major histocompability complex
                                                                                                                                                                                                                                                                                                                                                                                 Mutated forms (AAT41710-13) of the murine major histocompability complex interferon-stimulated response element (MMC IRSE) binding sequence (AAT41709), along with other 'competitor' DNAs (AA731714-16), were used in gel shift assays designed to determine whether mouse lymphocyte-specific interferon regulatory factor (LSIRF) (see also AAR99426) is a DNA binding protein. Mutant misRE mutant mt4 (AAT41713) competed well with wild-type MHC ISRE for binding to LSIRF protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Polymorphism, biallelic; human; forensic; paternity testing; disease; detection; phenotypic typing; characteristic; infection; hereditary; autoimmune disease; cancer; inflammation; drug; therapy; medicament; treatment; marker; primer; ss.
                                                                                                                                                             New genes for murine lymphocyte specific interferon regulatory factor used for modulation of lymphocyte activation and proliferation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New isolated mucleic acid segments from the human genome - used for determining polymorphic forms for use in e.g. forensics, paternity testing or phenotypic typing for disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
3.1%; Score 13.2; DB 1; Length 18;
Best Local Similarity 83.3%; Pred. No. 3e+02;
Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human biallelic polymorphic marker downstream primer #393.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 18 BP; 7 A; 2 C; 6 G; 3 T; 0 U; 0 Other;
Richardson CD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (WHED ) WHITEHEAD INST BIOMEDICAL RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 16; Page 197; 310pp; English.
                                                                                                                                                                                                                                                                                                Example 4; Page 41; 92pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2 GCCAGGAGTGAAACTGCG 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hudson T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GCTAGAAGTGAAACTGAG 18
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cc method for determining polymorphic forms in an individual for use in e.g. forensics, paternity testing or for phenotypic typing for diseases such as agammaglobulinemia, diabetes insipidus, Lesch-Nyham syndrome, muscular dystrophy, Wiskott-Aldrich syndrome, Fabry's disease, familial cypercholesterolemia, polycystic kidney disease, hereditary hereditary hasmorthosis, von Willebrand's disease, tuberous sclerosis, hereditary hasmorthagic telangiectasia, familial colonic polyposis, Bhlers-Danlos syndrome, osteogenesis implemention, cancer, diseases of the nervous syndrome, osteogenesis inflammation, cancer, diseases of the nervous system, infection by pathogenic microorganisms, and characteristics such as longevity, appearance (e.g. baldness, obseity), strength, speed, candurance, fertility, and susceptibility or receptivity to particular drugs or therapeutic treatments. The isolated polymorphic nucleic acid segments can also be used to produce medicaments for the treatment or prophylaxis of such diseases

Sequence 18 BP; 4 A; 6 C; 5 G; 3 T; 0 U; 0 Other;

DB 1; Length 18; Score 13.2; DB 1; Length 19 Pred. No. 3e+02; 0; Mismatches 3; Indels ., 344 CCGCCTCCTCTACAGCGA 361 3.1%; Query Match Best Local Similarity 83.3' 8

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Gaps ö

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AAX80491 RESULT

AAX80491;

AAX80491 standard; DNA; 18 BP.

(first entry)

26-AUG-1999

Human secreted protein yc2_1 probe.

Human, secreted protein, immunostimulator, nutrition, cytokine, cell proliferation, differentiation, immune stimulating, vaccine, suppression, haematopolesis regulation, tissue growth, activin, inhibin, chemotactic, chemokinetic, haemostatic, thrombolytic, anti-inflammatory; cadherin; tumour invasion suppressor; tumour inhibition; gene therapy; probe; hybridisation; ss. 

Homo sapiens. Synthetic

WO9932614-A1

01-JUL-1999

98WO-US027140. 18-DEC-1998; 97US-0068379P. 98US-00212843. 20-DEC-1997; 16-DEC-1998;

(GEMY ) GENETICS INST INC.

, Evans C; Wong GG, Clark HF; Collins-Racie LA, Steininger RJ, W Lavallie ER, Agostino MJ, Mccoy JM, Treacy M, Jacobs K, 1 Merberg D, Fechtel K;

WPI; 1999-395405/33.

New polynucleotides encoding secreted human proteins potentially useful e.g. immunostimulators.

Disclosure; Page 96; 99pp; English.

The present invention describes human secreted proteins obtained from human fetal brain, fetal kidney or adult blood cDNA libraries. The present sequence represents a probe for a human secreted protein. The human secreted proteins, and polynucleotides encoding them, are predicted human secreted proteins, and polynucleotides encoding them, are predicted

ö to have biological activities which would make them suitable for treating, preventing or ameliorating medical conditions in humans and animals, although no supporting data is given. Suggested activities include nutritional activity, cytokine and cell proliferation/differentiation activity, haematopoiesis requlating (e.g. as vaccines) or suppressing activity, haematopoiesis requlating activity, tissue growth haemacostatic and thrombolytic activity, receptor/ligand activity, anti-inflammatory activity, cadherin/tumour invasion suppressor activity, and tumour inhibition activity. The polynucleotides are also stated to be G-alpha-13; human; inhibitor; cancer; antisense compound; therapy; ss. Gaps ö Antisense compound inhibiting expression of human G-alpha-13 Query Match
3.1%; Score 13.2; DB 1; Length 18;
Best Local Similarity 83.3%; Pred. No. 3e+02;
Matches 15; Conservative 0; Mismatches 3; Indels Human G-alpha-13 antisense inhibitor ISIS# 20742. Sequence 18 BP; 2 A; 3 C; 8 G; 5 T; 0 U; 0 Other; 30 GGCTGGGACGAAGATGGC 47 1 Grcregaacearerresc 18 BP. 98US-00205860. 98US-00205860. AAZ31793 standard; DNA; 18 24-JAN-2000 (first entry) useful for gene therapy (ISIS-) ISIS PHARM INC. WPI; 1999-633376/54. Synthetic. Homo sapiens. 04-DEC-1998; 04-DEC-1998; US5981732-A. .99-NOV-1999. Cowsert LM; AAZ31793; RESULT 283 AAZ31793 ઠ g

ô This sequence represents an antisense inhibitor of the invention, and inhibits the expression of the human G-alpha-13 protein. The antisense compounds of the invention are of 8 to 30 nucleobases in length, that inhibits the expression of the human G-alpha-13. The antisense compound is useful for treating an animal, particularly humans, having or being prone to a disease or condition associated with the expression of G-alpha Gaps ö Score 13.2; DB 1; Length 18; Pred. No. 3e+02; 0; Mismatches 3; Indels Sequence 18 BP; 4 A; 6 C; 8 G; 0 T; 0 U; 0 Other; Claim 11; Col 38; 38pp; English. -13, such as cancer

103 CTGACCGCGACCGCAGCA 120 Query Match
Best Local Similarity 83.3%;
Matches 15; Conservative ઠે

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neutroparty; hardware damage; traumatic brain injury; peripheral neuropathy; Alzheimer's diesease; harkinson's diesease; parkinson's diesease; progressive Supranuclear Palsy; Olivopontocerebellar atrophy; Shy-Drager Syndrome; damannian parkinsoniam dementia complex; amyotrophic lateral sclerosis; memory impairment; neuropathy; ischemic stroke; acute brain injury; acute spinal cord injury; nervous system tumour; multiple sclerosis; eye disorder; pCR primer; se.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New isolated polypeptides, used for treating e.g. neurodegenerative disease or disorder, neuronal damage or neuronal disorder of the peripheral nervous system, the medulla or the spinal cord.
                                                                                 PCR primer NBNint.sense for neublastin neurotrophic factor cDNA.
                                                                                                     Neurotrophic factor; neublastin; neurodegenerative disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Johansen TE, Blom N, Hansen C;
                                                                                                                                                                                                                                                                                                                                      98US-0092229F.
98DK-00001048.
98US-0097774F.
98DK-00001265.
                  AAZ60571 standard; DNA; 18 BP.
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                                                             (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-171013/15.
                                                                                                                                                                                                                                                             WO200001815-A2.
                                                             05-MAY-2000
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                                        AAZ60571;
                                                                                                                                                                                                                                        Mus sp.
RESULT 284
           AAZ6057
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c factor designated neublastin. Neublastin is a member of the glial cell
c factor designated neublastin. Neublastin is a member of the glial cell
c factor-detaived neurotrophic factors sub-class of the transforming growth
c factor-beta superfamily of neurotrophic factors. Neublastin exhibits high
affinity for the GFR-alpha-RET receptor complex. The polypeptides can be
used for treating a neurodegenerative disease or disorder, cerebral
ischemic neuronal damage, traumatic brain injury, peripheral neuropathy.
Alzheimer's disease, Huntington's disease, Parkinson's disease, Parkinson
arrophy, Shy-Drager Syndrome, Guamanian parkinsonism dementia complex,
amyotrophic lateral solerosis, memory impairment, or a neuronal disorder
c of the peripheral nervous system, the medulla or the spinal cord. They
can also be used for treating various neuropathies. They can also be used
for treating ischemic stroke, acute brain injury, acute spinal cord
injury, nervous system tumours, multiple solerosis, exposure to
c neurotoxins, metabolic diseases such as diabetes or renal dysfunctions
and damage caused by infectious agents, or various disorders in the eye Claim 33; Page 32; 106pp; English.

Sequence 18 BP; 3 A; 8 C; 6 G; 1 T; 0 U; 0 Other;

3.1%; Score 13.2; DB 1; Length 18;

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Sequences AAZ49796-Z59835 represent antisense oligonuclectides targetted to the human Smad3 gene, which inhibit its expression. The antisense coligonuclectides were designed to target different regions of the human conjuguouclectides were designed to target different regions of the human smad3 mRA, and were analysed for their effect on Smad3 mRAA levels by canditative real-time PCR. The Smad proteins are a family of cytosolic proteins which are involved in TGF-beta superfamily signal transduction. On ligand binding, TGF-beta superfamily proteins (such as bone morphogenetic protein (BMP), activin and TGF-betas themselves) and proteins which then home or heterodimerise and translocate to the nucleus to activite rarget gene transcription. Smad3 translocate to the nucleus to activite respective of subgroup of Smad cannot factors, the pathway-restricted Smads, which are regulated by TGF-beta and activins. It can heterodimerise with Smad4 (196013787-A, AAY69622), the complex being able to activate TGF-beta conformed and cativins. It can heterodimerise with Smad4 canding the transcription. The oligonucleotides of the invention are useful for diagnosis, prevention and treatment of conditions associated with smad4 shades.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antisense inhibition of the human Smad3 gene, useful for diagnosing, preventing and treating conditions associated with Smad3 expression e.g.
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                                                                                                                                                                                                                                                                            Human Smad3 phosphorothioate antisense oligonucleotide, SEQ ID NO:9.
                                                                                                                                                                                                                                                                                                          Smad3, MADH3; hMAD3; JV15-2; TGF-beta signalling pathway;
transcription factor; expression inhibition; antisense therapy;
tumour formation; inflammation; antisense; ss.
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                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 18 BP; 1 A; 4 C; 12 G; 1 T; 0 U; 0 Other;
                  3
83.3%; Pred. No. 3e+02;
ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       331
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                                                      45 GGCCACCACTCAGAGGAG 62
                                                                                1 GGCCACGCTCCGACGAG 18
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Best Local Similarity 83.3
Matches 15; Conservative
                  15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2000-126072/11.
 Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Monia BP,
                                                                                                                                                                                                               AAZS9797;
                                                                                                                                          RESULT 285
                                                                                                                                                            AAZ5979
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The invention relates to novel human secreted proteins, the nucleic acids encoding them. The protein may exhibit cytokine, cell proliferation or call differentiation activity or may induce production of other cytokines in certain cell populations and may exhibit immune stimulating or immune suppressing activity, which is useful for the treatment of various immune deficiencies and disorders e.g. severe combined immunodeficiency (SCID), autoimmune disorders e.g. multiple sclerosis, systemic lupus erythematosus, rheumatoid arthritis, autoimmune pulmonary inflammation. The proteins are also useful in the treatment of diseases and disorders including tissue, skin and organ transplantation and in graft-versue-host diseases (GVBD), in the induction of tumour immunity, myeloid or lymphoid cell deficiencies, wound healing and tissue repair, in the treatment of burns, incisions and ulcers, as well as in treatment of periodontal disease, osteoporosis or osteoarthritis, mediated by inflammatory processes, diseases (Hunington's disease, amylotrophic lateral selections and SNy-Drager syndrome, infections, inflammatory bowel disease, ulcers, bone regeneration. The protein, having activin- or inhibin-related activities is useful as a contraceptive based on the ability of inhibins male mammals. The proteins and nucleic acids are also spermatogenesis in male mammals. The proteins and nucleic acids are also useful as food supplements. The present sequence is an oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Clark H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Secreted human proteins, useful as vaccine for treating various diseases such as autoimmune disorders (e.g. multiple sclerosis), and nervous system disorders (e.g. stroke).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nundari section de protective, antiantimitation, antimitatobial, vulnerary, nontropic, neuroprotective, antiantimitation, antimitatobial, vulnerary, cytostatic, antidiabetic; virucide, antinifertility, anticonvulsant; vasotropic, antiparkinsonian; immunoetimulant; dermatological; probe; anticheumatic; antitumor; antiulcer; osteopathic; tranquiliser; cerebroprotective; cytokine; cell proliferation; cell differentiation; immune deficiency; sCID; tumour; autoimmune disorder; multiple sclerosis; rheumatoid arthritis; autoimmune disorder; multiple sclerosis; neumatoid arthritis; periodotal disease; osteoporosis; ost
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Evans C;
Wong GG,
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Agostino MJ, Steininger RJ, Spaulding V,
Merberg D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 protein; ss; antiinflammatory;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human secreted protein yc2_1 probe.
                                                                                                                        AAS59326 standard; DNA; 18 BP.
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04-DEC-2000; 2000US-00729674.
                                                                                                                                                                                                                                                                                                                                                                                16-JAN-2002 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-639363/73.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        secreted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200175068-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11-OCT-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Treacy M,
Fechtel K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Jacobs K,
Treacy M,
                                                                                                                                                                                                                                                       AAS59326;
RESULT 286
AASS9326
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to an SCR (Sequence Characterised amplified Regions) primer for distinguishing Korean beef meat from milk cow meat. The SCR primer distinguishes between the two quickly and accurately. The current sequence represents an SCR primer for distinguishing between beef
probe used to detect the nucleic acids of the invention and where an N residue is present at position 2 this is a biotinylated phosphoroamidite residue
                                                                                                                                                                                                                                                                                                                      SCR; sequence characterised amplified regions; beef; cow; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3.1%; Score 13.2; DB 1; Length 18; 33.3%; Pred. No. 3e+02; ve 0; Mismatches 3; Indels
                                                                                Length 18;
                                                                                                        3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Yoon DH;
                                                                                                                                                                                                                                                                                               SCR primer 1 for distinguishing between beef types.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 18 BP; 3 A; 6 C; 7 G; 2 T; 0 U; 0 Other;
                                                                             Score 13.2; DB 1;
Pred. No. 3e+02;
0; Mismatches 3;
                                                        BP, 2 A, 3 C, 8 G, 5 T, 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Scr primer for distinguishing korean beef meat.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hong YH, Jung IJ, Kim HB, Kim HS, Kim TH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (RURA-) RURAL DEV ADMINISTRATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 5; 6pp; Korean.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   302 CCTGAGCCCCGGGGACCG 319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CCAGAGCCTCGGGGACTG 18
                                                                                                                                 30 GGCTGGGACGAAGATGGC 47
                                                                                                                                                 1 GTCTGGGACGATGTTGGC 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                        99KR-00033412.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 83.3%;
Matches 15; Conservative
                                                                                Query Match 3.1%;
Best Local Similarity 83.3%;
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABT11899 standard; DNA; 18
                                                                                                                                                                                                                       ABL53448 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-495317/54.
                                                                                                                                                                                                                                                                                                                                                                                     KR2001017747-A.
                                                                                                                                                                                                                                                                                                                                                              Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              13-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                        13-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-DEC-2002
                                                                                                                                                                                                                                                                       31-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                              05-MAR-2001.
                                                         Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABT11899;
                                                                                                                                                                                                                                                ABL53448;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 288
                                                                                                                                                                                               RESULT 287
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                                                                                                                                                                                                            ABL53448
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neuroprotective; haemostatic; thrombolytic; anti-inflammatory; phosphoarmidite; ss.

97US-0126425P. 97US-0067454P. 97US-0070345P. 98US-0070643P. 98US-0071345P. 98US-0071344P.

02-JAN-1998 07-JAN-1998

08-JAN-1998 13-JAN-1998 22-JAN-1998 30-JAN-1998

20-DEC-1997

2000US-00729674

04-DEC-2000;

08-NOV-2001

US2001039335-A1.

Synthetic

98US-0075038P

18-FEB-1998; 23-NOV-1998;

2000US-00539330

MCCOY J M.
LAVALLIE E R.
COLLINS-RACIE L

JACOBS

(JACO/)

EVANS C. MERBERG D. TREACY M. AGOSTINO M J. STEININGER R J

(MCCO/) MCCOY J M.
(LAVA/) LAVALLIE E I
(COLL/) COLLINS-RAC!
SVAN/) EVANS C.
(MERB/) MERBERG D.
(TREA/) TREACY M.
(AGOS/) AGOSTINO M.
(STEL/) STEININGER I
(SPRU/) SPAULDING V
(MONG/) WONG G G.
(CLAR/) CLARK H.

SPAULDING V.

98US-0073095P

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The invention relates to a truncated neublastin polypeptide comprising an amino acid terminus that lacks one or more amino-terminal amino acids of a mature neublastin polypeptide. The polypeptides and mucleic acids are useful for treating neurodegenerative disorders such as ischemic neuronal damage, traumatic brain injury, peripheral neuropathy, neuropathic pain, Alzheimer's disease, Huntington's disease, Parkinson's disease, amyotrophic lateral sclerosis, memory impairment, diabetes, renal diseases, or glaucoma by moderating metabolism, growth, differentiation or survival of a nerve or neuronal cell. This polynucleotide sequence is a neublastin PCR primer of the invention
                                     Nootropic; neuroprotective; antiparkinsonian; anticonvulsant; analgesic; tranquiliser; antidiabetic; ophthalmological; neurodegenerative disorder; neublastin; ischemic neuronal damage; traumatic brain injury; diabetes; peripheral neuropathy; neuropathic pain, Alzheimer's disease; glaucoma; Huntington's disease; Parkinson's disease; amyotrophic lateral sclerosis; memory impairment; renal disease; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New truncated neublastin polypeptides lacking one or more amino-terminal amino acids of a mature neublastin polypeptide useful for treating neurodegenerative disorders, e.g. peripheral neuropathy, neuropathic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18 BP; 3 A; 8 C; 6 G; 1 T; 0 U; 0 Other;
Neublastin DNA related PCR primer SEQ ID No
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rossomando A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Page 44; 138pp; English.
                                                                                                                                                                                                                                                                                                                12-MAR-2002; 2002WO-EP002691
                                                                                                                                                                                                                                                                                                                                                          12-MAR-2001; 2001US-00804615
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sah DWY, Johansen TE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-713515/77.
                                                                                                                                                                                                                                                                                                                                                                                              (BIOJ ) BIOGEN INC.
(NSGE-) NS GENE AS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              pain, brain injury.
                                                                                                                                                                                                                                  WO200272826-A2.
                                                                                                                                                                                        Unidentified.
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The invention relates to isolated polymucleotides (ABA90876-ABA90968 and ABA9080) and encoded proteins (ABB55698-ABB55800), especially conjunciocidectides SEQ ID NO 12 (ABA908767) and SEQ ID NO 19 (ABA90885) and SEQ ID NO 19 (ABA90885) and SEQ ID NO 20 (ABB55707) contained in clones bd306-7 and yb8-1 respectively and the clones bd306-7 and yb8-1 respectively and the clones bd306-7 and yb8-1 respectively and the clones bd306-7 and yb8-1 are deposited with the American Type Culture Collection (ArCO) with accession number 98599. The polymucleotides and encoded polypeptides have cycostatic, anti-inflammatory, immunomodulator, vulnerary, chrombolytic and anti-inflammatory activity and acting as cytokine modulators, cand anti-inflammatory activity and acting as cytokine modulators, cand anti-inflammatory activity and acting as cytokine modulators, chamatopoiesis regulators, tissue growth modulators and/or cadherin charapies, particularly for preventing, treating or amelicrating any of the following diseases: immune deficiency and disorders; e.g. bacterial or fungal infections, autoimmune disorders, categorousis or cytomatosus or graft-versus-host disoases; myeloid or lymphoid cell deficiencies; wound, burns, incisions and ulcers, osteoporosis or cytomathies, e.g. Alzheimer's, Parkinson's disease, Huntington's characterial selection or stroke, inflammations, shock, sepais
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New secreted proteins and encoding polynucleotides, useful in gene therapies, particularly for preventing or treating autoimmune disorders, cancer, graft-versus-host disease, wound, osteoporosis, stroke or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Collins-Racie LA, Evans C;
Steininger RJ, Spaulding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 329; 349pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lavallie ER,
1, Agostino MJ,
Fechtel K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       D, Treacy M,
Clark H, Fe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mccoy JM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-040725/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        inflammations.
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Wong GG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Jacobs K,
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Gaps ò

Query Match 3.1%; Score 13.2; DB 1; Length 18; Best Local Similarity 83.3%; Pred. No. 3e+02; Matches 15; Conservative 0; Mismatches 3; Indels

GGCCACCACTCAGAGGAG 62 GGCCACCGCTCCGACGAG 18 or systemic inflammatory response syndrome, ischaemia-reperfusion injury

Human; clone bd306-7; clone yb8-1; ATCC number 98599; gene therapy; immune disorder; bacterial infection; fungal infection; cancer; tumour; autoimmune disorder; spetemic lupus erythematosus; wound; ulcer; inhibin; osteoporosis; osteoarthritis; nervous system disorder; neuropathy; Alzheimer's disease; Parkinson's disease; Huntington's disease; activin; hamemophilia, cardidar infarction; stroke; sepsis; arthritis, vulnerary; ischaemia-reperfusion injury; inflammatory bowel disease; chemctactic; crohn's disease; cytostatic; anti-inflammatory; immunomodulator;

Biotinylated oligonucleotide SEQ ID NO 213.

(first entry)

14-FEB-2002

ABA90995;

ABA90995 standard; DNA; 18

289

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**ABA9099** RESULT

Endoplasmic reticulum stress competence control element SEQ ID NO:11.

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AAZ25638 standard; DNA; 19

AAZ25638,

23-DEC-1999

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AAZ25638;

Endoplasmic reticulum; ER; stress competence; control element; inhibition; growth; apoptosis; cancer; autoimmune disease; cystic fibrosis; ds.

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The present sequence represents a specifically claimed oligonucleotide PCR primer. The oligonucleotide can be used for polymerase chain reaction (PCR) amplification of DNA, specifically regions of specific genes that are conserved among mammalian species, i.e. pairs of oligonucleotides from the present specification represent universal mammalian sequence-tagged site (UM-STS) primers. The primers are used to develop genomic maps, to isolate clones from libraries, to make cross-species comparisons and to develop additional genetic markers. UM-STS allow genomic comparisons to be made between more species
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New oligonucleotide primers amplifying gene regions conserved among mammals - useful for developing genomic maps, isolating clones and making cross-species comparisons.
endotoxin lethality, arthritis, inflammatory bowel disease or Crohn's disease; or tumours or cancers, pemphigus vulgaris or pemphigus foliaceus. The present sequence is that of a blotinylated oligonuclectide with a phosphoaramidite residue, useful to the invention
                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer; polymerase chain reaction; amplification; UM-STS; universal mammalian sequence tagged site; genomic map; clone; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                             Interleukin 2 receptor PCR primer for universal mammalian STS's.
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                                                                                                                                DB 1; Length 18;
                                                                                                                                                                   3; Indels
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                                                                                             Sequence 18 BP; 2 A; 3 C; 8 G; 5 T; 0 U; 0 Other;
                                                                                                                            3.1%; Score 13.2; DB 1
Local Similarity 93.3%; Pred. No. 3e+02;
es 15; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Yuzbasiyan-Gurkan V;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 10; 26pp; English.
                                                                                                                                                                                                                                       18
                                                                                                                                                                                                      30 GGCTGGGACGAAGATGGC 47
                                                                                                                                                                                                                                                                                                                                    AAV01209 standard; DNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      96US-0012061P.
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(UNMS ) UNIV MICHIGAN STATE.
                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1997-435083/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18-FEB-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                              23-MAR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9731012-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                         AAV01209;
                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                     RESULT 290
                                                                                                                                                                       Matches
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New control element for stress competence of endoplasmic reticulum - useful for inhibition of growth and induction of apoptosis in cancer

98JP-00052453 98JP-00052453

04-MAR-1998; 04-MAR-1998;

JP11243959-A.

Gallus sp.

(HSPK-) HSP KENKYUSHO KK. WPI; 1999-603708/52.

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The present invention specifically claims an element shown by: (A) a 19 bp base sequence, CCAATNNNNN NNNNCCACG (ERSB); or (B) a modified base sequence having replaced 1.3 bases with the other base(s), which induces transcription with stress on endoplasmic reticulum used for stress competence of endoplasmic reticulum. Also described are: (1) a DNA having transcription inducing activity with stress on endoplasmic reticulum containing the above mentioned element, optionally further containing a promoter DNA, and (2) a vector containing the element can be used for the inhibition of growth and induction DNA. The element can be used for the inhibition of growth and induction dispasses and cystic fibrosis by inhibition of symptoms of canter colls, and improvement of symptoms of canter calls, and improvement of symptoms of canter calls.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3.1%; Score 13.2; DB 1; Length 19; Similarity 83.3%; Pred. No. 3.46+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19 BP; 4 A; 9 C; 5 G; 1 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GRP94 promoter ERSE3-like sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Fig 3; 25pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    150 GAGGCCGGCTTCGACTGG 167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP.
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AAA28576 standard; DNA; 19
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        present invention
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29-AUG-2000
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Best Local S:
Matches 15
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Gape

350 GCTCTACAGCGACTTCCT 367

18

gcrcracagagarccr

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An endoplasmic reticulum stress transcription factor (bZIP) capable of regulating transcription inducing activity exhibited by an element (ERSE) can be used in a method for controlling expression of an endoplasmic reticulum chaperone. The method comprises expressing bZIP. The method can be used for expression of a foreign protein by positively regulating expression of an endoplasmic reticulum chaperone gene. bZIP is useful for controlling the expression of endoplasmic reticulum chaperone either positively or negatively in cells and therefore is useful for treatment or prophylaxis of encers and endoplasmic reticulum chaperone either or prophylaxis of encers. bZIP also maintains the correct conformation diseases, wounds and ulcers. bZIP also maintains the correct conformation of the endoplasmic reticulum chaperone and thereby increases the expression of a foreign protein. This sequence taken from the glucose regulating protein (GRP) promoter GRP94 contains an BRSE like sequence. (Updated on 15-SEP-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                New endoplasmic reticulum stress transcription factor (known as bZIP) for controlling expression of endoplasmic reticulum chaperone, useful for treating cancers, arteriosclerosis, cystic fibrosis, ischemic diseases,
chicken; gene expression; GRP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 19 BP; 4 A; 9 C; 5 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                       Yura T;
                                                                                                                                                                                                                                                                                                                       Mori K, Yanagi H,
  ulcer; gene therapy; recombinant gene; c. glucose regulated protein; promoter; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Fig 3; 157pp; English.
                                                                                                                                                                                 99WO-JP006305
                                                                                                                                                                                                                     98JP-00324227
                                                                                                                                                                                                                                                                                 (HSPR-) HSP RES INST INC.
                                                                                                                                                                                                                                                                                                                         Yoshida H,
                                                                                                                                                                                                                                                                                                                                                                WPI; 2000-387736/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                 treating cancers, wounds and ulcers.
                                                                                                    WO200029429-A2.
                                                             Gallus gallus
                                                                                                                                                                                                                         13-NOV-1998;
                                                                                                                                                                                                                                            09-JUN-1999;
                                                                                                                                                                                   12-NOV-1999;
                                                                                                                                          5-MAY-2000.
                                                                                                                                                                                                                                                                                                                           Haze K,
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Nucleic acids encoding human kinase polypeptides, useful for preventing diagnosing and/or treating e.g. cancer, immune, cardiovascular and neuronal-associated diseases, and microbial infections.

Sudarsanam S, Martinez R;

Manning G,

Whyte D, Clary D;

Plowman GD, Flanagan P,

WPI; 2001-343950/36.

22-NOV-2000; 2000WO-US032085.

99US-0167482P

24-NOV-1999;

(SUGE-) SUGEN INC.

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Canine distemper virus H gene PCR primer RH-3.
                                                         35 GGACGAAGATGGCCACCA 52
                                                                                             18 GGCCAAAGATGGCCTCCA 1
                                                                                                                                                                                                           絽
                                                                                                                                                                                                           AAF86572 standard; DNA; 19
                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                             Canine distemper virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        JP2000350587-A.
                                                                                                                                                                                                                                                                                               12-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-DEC-2000.
                                                                                                                                                                                                                                                        AAF86572;
Best Loca
Matches
                                                                                                                                                                    RESULT 294
                                                                                                                                                                                             4AF86572/
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                                                                                                                                                                                                                                                                                                                                                                                                                               Human; protein kinase; PTK; STK; cancer; cardiovascular disease; SNP; metabolic disorder; immune related disease; neurological disorder; neurodegenerative disorder; inflammatory disorder; infectious disease; reproductive disorder; gene therapy; single nucleotide polymorphism; ds
                                                                    Gaps
                                                                  ö
                    3.1%; Score 13.2; DB 1; Length 19; 83.3%; Pred. No. 3.4e+02; tive 0; Mismatches 3; Indels
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SNP containing protein kinase DNA sequence #54.

WO200138503-A2

31-MAY-2001

Homo sapiens

(first entry)

12-SEP-2001

AAS06885;

150 GAGGCCGGCTTCGACTGG 167

15; Conservative

Query Match Best Local Similarity

Best Loca Matches

credcececercarred 1

18

g ò

AAS06885 standard; DNA; 19

AAS06885/

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AASO6832-AASO6897 represent part of a polymucleotide sequence encoding for novel human protein kinases where a single nucleotide polymorphism (SNP) has been identified. The SNP occurs at the last position of the present sequence. The sequences are described relating to the invention of novel human protein kinases #1-57 (AMU03501-AAU03557). The novel protein kinases (ATK and 1-57 (AMU03501-AAU03557). The novel control of the protein kinases (ATK and STK) families. The polymorlootides serine/threonine kinase (ATK and STK) families. The polymorlootides prevention, diagnosis and treatment of diseases associated with imappropriate kinase expression. For example, they may be used to treat cancers (especially cancers of haematopoietic origin), cardiovascular disease (e.g. atherosclarosis), membolic disorders (e.g. diabetes), immune related diseases (e.g. rheumatoid arthritis), neurological immune related diseases (e.g. rheumatoid arthritis), neurological protein kinase (e.g. disease), inflammatory disorders (e.g. asthma), infectious disease (e.g. HIV) and reproductive disorders (e.g. asthma), infectious disease (e.g. HIV) and reproductive disorders (e.g. infertility).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene therapy and as DNA probes in diagnostic assays. The protein Kinase polypeptides may be used as antigens in the production of antibodies against the protein kinases and in assays to identify modulators of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Canine; H gene; antiviral; gene therapy; distemper; PCR primer; ss.
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               / Match
Jocal Similarity 83.3%; Pred. No. 3.4e+02;
los 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 19 BP; 2 A; 5 C; 6 G; 5 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 protein kinase expression and activity
                                                                                                                                                                                                                                                                          Example 8B; Page 333; 433pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        99JP-00165598
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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(KYOR-) KYORITSU SHOJI KK.

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The invention relates to a method for determining whether an individual is likely to be susceptible to malignant melanoma, and determining the genetic basis for the melanoma in an individual. The method involves screening the genome of the individual for the presence or absence of one or more polymorphic variants of the XRCC3 gene. Sequences AA447412-420 represent PCR primers used in a genotyping assay of a candidate DNA repair gene XPD
                                                                                                                                       The present invention relates to the H gene derived from canine distemper virus (see AAF6557). The H gene sequence can be used in the prevention, treatment and detection of mammalian distemper, particularly canine distemper virus (CDV). The present sequence is a PCR primer, which was used in the present invention
                                                         H gene, used for treating, preventing and detecting mammalian distemper, particularly canine distemper viruses.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Determining the susceptibility of an individual to malignant melanoma, involves screening the genome of the individual for the presence or absence of one or more polymorphic variants of the XRCC3 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         XRCC3; XPF; melanoma; genotyping; DNA repair gene; XPD; PCR primer;
                                                                                                                                                                                                                                                                      3.1%; Score 13.2; DB 1; Length 19; 83.3%; Pred. No. 3.4e+02; tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 19 BP; 6 A; 4 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                         Sequence 19 BP; 4 A; 6 C; 6 G; 3 T; 0 U; 0 Other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Welsh K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Wojnarowska F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          XPD gene exon 23 amplifying primer.
                                                                                                             Example 2; Page 6; 18pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example; Page 14; 35pp; English.
                                                                                                                                                                                                                                                                                                                                          293 GGTGAAGGACCTGAGCCC 310
                                                                                                                                                                                                                                                                                                                                                                          N
                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22-FEB-2000; 2000GB-0004193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ISIS-) ISIS INNOVATION LTD
                                                                                                                                                                                                                                                                                                                                                                           GCTGGAGTACCTGAGCCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAH47419 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 83.3
Matches 15, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Haldar N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-557711/62.
                           WPI; 2001-268280/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            polymorphism; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200162964-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-NOV-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAH47419;
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                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 295
                                                                                                                                                                                                                                                                                                                                                                                                                                          AAH47419
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The present invention describes a method of arraying genome clones. The method comprises: (a) clones of the genomic libraries contained in method comprises numbered for discrimination are mixed in each of the multiwell plates; (b) a primer designed based on the chromosome marker completed protected from the resultant corresponding to the marker is detected from the resultant corresponding to the marker is detected from the resultant corresponding to the marker is detected from the resultant corresponding to the marker is detected from the marking is changed so that the same discrimination Nos. of the multiwell cof the maximum in the specified discrimination Nos. to array the multiwell cof the maximum in the specified discrimination Nos. to array the multiwell corresponding to that the same discrimination Nos. succeed to discrimination Nos. are mixed respectively in each wells of longitudinal corresponding to the mixed clones are cultured and the call lateral directions; (f) the mixed clones are cultured and the created from the amplified products; (h) the clones in the multiwell corresponding from the detected result; and (i) the clones are constituted as the positions on the chromosome and arrayed. The crons in the multiwell corresponds for human chromosome 126-35 DNA, and ABM45323 to ABM45634 crepresent PCR primers for human chromosome 21q22.1, which are corrected in the present invention
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ö
                                                                                                                                                                                                                                                                                                     Human; chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome;
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                                                                                                                                                                                                                                                                   Human chromosome 1p36-35 PCR primer SEQ ID NO:1028.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 19 BP; 5 A; 7 C; 4 G; 3 T; 0 U; 0 Other;
   3,
   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 4; Page 25; 528pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GTGCTGGCGGCGGACGAC 338
   ..
                                       69
                                                                     AATCAGAGGAGACGCTGC 19
                                                                                                                                                            ABL43984 standard; DNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12-MAR-2001; 2001JP-00068285.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-MAR-2000; 2000JP-00066716.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    83.3%;
                                       52 ACTCAGAGGAGTCTCTGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (RIKA ) RIKAGAKU KENKYUSHO.
(GENO-) GENOTEX YG.
                                                                                                                                                                                                                                  11-APR-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity 83.3
nes 15; Conservative
     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Arraying genome clones.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-144136/19
                                                                                                                                                                                                                                                                                                                             PCR primer; sa
                                                                                                                                                                                                                                                                                                                                                                                                 JP2001321190-A
                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                   20-NOV-2001.
       15;
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                                                                                                                                                                                                  ABL43984;
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       Matches
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Gaps

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13

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RESULT 297

Length 19;

3.1%; Score 13.2; DB 1; 83.3%; Pred. No. 3.4e+02;

Query Match Best Local Similarity

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Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                           Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; SEQ ID NO 12494; 872pp; English.
                                                                                                                                                                                                                                                                                                          Li Y, Sandrasagra A,
Tang L, Shahabuddin
          ABZ97252 standard; DNA; 19 BP
                                                                                                                                                                                                                                                                24-APR-2001; 2001US-0286137P.
                                                                                                                                                                                                                                           23-APR-2002; 2002WO-US013135.
                                                                                                                                                                                                                                                                                    EPIG-) EPIGENESIS PHARM INC
                                                                          Human nucleic acid sequence.
                                                    17-0CT-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                           WPI; 2003-229219/22.
                                                                                                                                                                                                WO200285308-A2.
                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                     31-OCT-2002.
                                                                                                                                                                                                                                                                                                          Nyce JW, 1
Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                 ubiquinone
                               ABZ97252;
ABZ97252
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Katz E, Pabalan J, Aguilar D; S;

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' or second active associated with lung and/or nasal airway dysfunctions and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory steroid and ubiquinone. A composition may have a immunosuppressive, and cytostatic activity. The composition may have a preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an autioniflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of, or receptor, producing bronchodilation, increasing levels of ubiquinone or lung suffactant in a subject, for reating bronchodilation, increasing levels of ubiquinone or lung inflammation, lung allergies, or a respiratory disease or condition.

Note: The sequence data for this patent is not represented in the printed of the thin wind interval to the printed of the thin increasing format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences

Sequence 19 BP; 2 A; 10 C; 3 G; 4 T; 0 U; 0 Other;

Gaps 3.1%; Score 13.2; DB 1; Length 19; 83.3%; Pred. No. 3.4e+02; ve 0; Mismatches 3; Indels 83.3%; 15; Conservative Similarity Local Best Loca Matches

ACCACGCCGGCTTCTCT

RESULT 298

337 ACCAGGCCGGCTGCTCT 354

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antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; admissiones gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
                                                                                                                                                                                                                                                                                                          antisense; lung dysfunction; nasal airway dysfunction;
                                                                                                                                                                                                                                           Human IL4-R oligonucleotide sequence.
                                        ABZ97333 standard; DNA; 19
                                                                                                                                                                         (first entry)
                                                                                                                                                                             17-OCT-2003
                                                                                                          ABZ97333;
ABZ97333

XXX XXX ABZ9

XXX XXX BUM

XXX XXX BUM

XXX XXX BUM

XXX ABZ9

XXX ABZ9

XXX ABZ9

XXX BUM

XXX ABZ9

XXX BUM

XXX BUM

XXX ABZ9

XXX BUM

XXX BUM

XXX ABZ9

XXX BUM

XXX ABZ9

XXX ABZ9
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Katz E, Pabalan J, Aguilar D; Li Y, Sandrasagra A, Ka , Tang L, Shahabuddin S; 23-APR-2002; 2002WO-US013135. 24-APR-2001; 2001US-0286137P. (EPIG-) EPIGENESIS PHARM INC Nyce JW,

W0200285308-A2 Homo sapiens.

31-OCT-2002.

Miller S,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 12575; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonuclectide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nuclectides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entitlenamatory steroid and ubiquinone. A composition of the invention has antiinflammatory steroid and ubiquinone. A composition of the invention composition antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing levels of adenosine or receptor, producing bronchodilation, increaging levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchodilation, contracting bronchodilation, increaging levels of ubiquinone or lung inflammatory than all all producing all and all producing bronchodilation, increaging levels of ubiquinone or lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO cut if the printed and for this patences.

Sequence 19 BP; 2 A; 10 C; 3 G; 4 T; 0 U; 0 Other;

Gaps ö Score 13.2; DB 1; Length 19; Pred. No. 3.4e+02; 0; Mismatches 3; Indels ; 0 Query Match
Best Local Similarity 83.3%;
Matches 15; Conservative C

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354 13 337 ACCAGGGCCGGCTGCTCT ACCACGCCCGGCTTCTCT

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RESULT 299

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25-MAR-2003
07-JUL-1992
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stem_loop
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ID AAQ225
XX
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    THXSXEXBXBXB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ò
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a novel isolated double stranded RNA oligomucleotide about 19 to about 25 ribonucleotides in length or its equivalent. One strand of the oligomucleotide comprises the same nucleotide sequence as a region of a hepatitis C virus (HCV) target RNA polymucleotide sequence required for hepatitis C virus infection, replication or pathogenesis in vitro or in vivo in a host cell. The oligomucleotide of the invention demonstrates virucide activity and may be useful for preparing a composition or vaccine for treating or preventing hepatitis C virus, as well as during gene therapy procedures. The current sequence is that of the anti-HCV agent LZ129 mutant RNA of the invention which contains a C3G mutation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                     /*tag= a
/note= "Wild-type cytosine substituted for guanine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New double stranded RNA oligonucleotide, useful for preparing a composition for treating or preventing hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ò
                                                                                                                                HCV infection; replication; pathogenesis; virucide; vaccine;
gene therapy; ds; anti-HCV; agent LZ129; mutant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3.1%; Score 13.2; DB 1; Length 19; 72.2%; Pred. No. 3.4e+02; tive 2; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19 BP; 3 A; 6 C; 7 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Zhao G, Lu J, Glass JI, Martinez A, Yang Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Anti-HCV agent LZ129 mutant RNA - G4C.
                                                                                                      Anti-HCV agent LZ129 mutant RNA - C3G.
                                                                                                                                                                                                          Location/Qualifiers misc_difference 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 2; Page 155; 173pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   256 CGGCCACGGTGCACCTGG 273
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 CGGGCACGAUGCAUCUGG 18
              ADD00872 standard; RNA; 19 BP.
                                                                                                                                                                                                                                                                                                                                                                                    17-AUG-2001; 2001US-0313076P.
20-DEC-2001; 2001US-0344116P.
01-FEB-2002; 2002US-0353750P.
                                                                                                                                                                                                                                                                                                                                                          16-AUG-2002; 2002WO-US021843.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADD00871 standard; RNA; 19
                                                                        (first entry)
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Best Local Similarity 72.2%
Matches 13, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-268345/26.
                                                                                                                                                                             Synthetic.
Hepatitis C virus.
                                                                                                                                                                                                                                                                                                WO2003016572-A1
                                                                         01-JAN-2004
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                                                                                                                                                                                                                                                                                                                             27-FEB-2003
                                            ADD00872;
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ADD00872
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The invention relates to a novel isolated double stranded RNA oligonucleotide about 19 to about 25 ribonucleotides in length or its equivalent. One strand of the oligonucleotide comprises the same nucleotide sequence as a region of a hepatitis C virus (HCV) target RNA polynucleotide sequence required for hepatitis C virus infection, replication or pathogenesis in vitro or in vivo in a host cell. The oligonucleotide of the invention demonstrates virucide activity and may be useful for preparing a composition or vaccine for treating or preventing hepatitis C virus, as well as during gene therapy procedures. The current sequence is that of the anni-HCV agent LZ129 mutant RNA of the invention which contains a G4C mutation.
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                                                                                                                                                                                                                                                                                 /*tag= a
/note= "Wild-type guanine substituted for cytosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New double stranded RNA oligonucleotide, useful for preparing a composition for treating or preventing hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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HCV infection; replication; pathogenesis; virucide; vaccine; gene therapy; ds; anti-HCV; agent LZ129; mutant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.1%; Score 13.2; DB 1; Length 19; 72.2%; Pred. No. 3.4e+02; Live 2; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19 BP; 3 A; 8 C; 5 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Zhao G, Lu J, Glass JI, Martinez A, Yang Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 2; Page 155; 173pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                          Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 CGCCCACGAUGCAUCUGG 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAQ22593 standard; RNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17-AUG-2001; 2001US-0313076P.
20-DEC-2001; 2001US-0344116P.
01-FEB-2002; 2002US-0353750P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-AUG-2002; 2002WO-US021843
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 72.2
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EGS; Viral diseases; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ELIL ) LILLY & CO ELI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-268345/26.
                                                                                                                                                                                                                                            misc_difference 19
                                                                                                     Synthetic.
Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                   WO2003016572-A1.
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/*tag= a
/note= "forms loop with substrate"
9. .15

/*tag= b /label= EGS

misc_RNA

WO9203566-A

Guerrierta CL;

Forster AC,

Altman S,

UYYA ) UNIV YALE. 17-AUG-1990;

4PI; 1992-096909/12

91WO-US005808 90US-00568834 Disclosure; Fig 2d; 34pp; English

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To develop antisense oligos, the test system employed mouse NIH 3T3 cells stably transfected with an internally deleted construct of the human gene for the pro alpha 1(1) chains of type I procollagen COLLAI). A series of modified oligos were synthesised using a region at the 3' end of excn 1 and the first two nucleotides of intron 1 of the exogenous (human) gene as a target. This sequence is given in AAQ66555 which corresp. to bps 198 -225 if the adenine at the start of transcription is counted as poon. The corresp, which corresp to bps 169-195. The antisense oligos are given in AAQ66557, which corresp to bps 169-195. The antisense oligos are given in AAQ66557. Geoficia. The antisense oligos in metant or normal collagen genes. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                 Anti:sense oligo:nucleotide(s) against mutant or native collagen genes for inhibiting collagen expression, e.g for treating osteoarthritis, liver cirrhosis, excessive scarring etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human type I procollagen (COLIA1) pro alpha 1 chain antisense oligonucleotide AS9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 0 A; 7 C; 5 G; 8 T; 0 U; 0 Other;
Procollagen; antisense oligo; inhibition; ss
                                                                                                                                                                                                                                                                                                                                Nugent P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Procollagen; antisense oligo; inhibition;
                                                                                                                                                                                                                                                                                                                              Baserga R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 5; Page 24; 55pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26 CGAGGGCTGGGACGAAGA 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAO66602 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                   (UYJE-) UNIV JEFFERSON THOMAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20 CGAGGGCCAAGACGAAGA
                                                                                                                                                                                                                                      92US-00973332
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                              WPI; 1994-183496/22.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-MAR-2003
10-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-MAY-1994
                                                                                                                                                                                                                                      09-NOV-1992;
                                                                                               WO9411494-A1
                                                                                                                                            26-MAY-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ66602;
                                                  Synthetic.
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AAO66602/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ઠ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The sequence is designed to bind to a truncated deriv. of ATI (McClain, C al., Science 238, 527-530 (1987), so targetting cleavage of this c aubstrate by Rhase P. ATI comparises the acceptor stem, the T-stem and cloop, and the 31 terminal NCCA nucleotides (nt) of the tRNA PHE gene. The cloop, and the 31 terminal NCCA nucleotides (nt) of the tRNA PHE gene. The cerulting linear DNA was trains. cribed in vitro with SP6 polymerase, transcription yielding a short 5' leader sequence, and an extra 3'C restriction site. A 51 nt deriv., pATI (see AAQ2289) lacking residues 25 of ATI was also prepd,, and used to create a truncated substrate concered by E. coli RNAse P or M. RNA under conditions where pATI was not cleaved by E. coli RNAse P or M. RNA under conditions where pATI was cleaved efficiently. However if the cleaved EGS sequence (shown here) was cleaved efficiently. However if the cleaved EGS sequence (shown here) was cleaved efficiently. However if the cleaved EGS sequence (shown here) added to the mixt., cleavage occured as normal. This led to the design of cased that he cleaved EGS sequence (shown here) cracet sequence) and a 3' NCCA terminal, (N = a purine). Compsns. context the presence of viral RNAs themselves. RNAse P based therapy may be used to deliver engineered sequences into the haematopoietic cells of patients with e.g. HIV, HILV-I and various retroviral induced leukaemias. (Updated on 25-MAR-2003 to correct PA field.) (Updated on 25-MAR-2003 to correct PA field.) (Updated on 25-MAR-2003 to correct PA field.)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Compsn. for targeting RNA sequence for cleavage by RN ase P - comprises external guide sequence including 3-NCCA and complementary nucleotide sequences, for treating viral diseases.
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Length 20; 3; Indels

Score 13.2; DB 1; Pred. No. 3.8e+02; 0; Mismatches 3;

Query Match
Best Local Similarity 83.3%;
Matches 15; Conservative

243 TGCTTCCCGGGCTCGGCC 260

19 IGGIGCCCGGACICGGCC

g

N

Seguence 20 BP; 3 A; 8 C; 8 G; 0 T; 1 U; 0 Other;

Human type I procollagen (COLIAI) pro alpha 1 chain antisense oligonucleotide AS8.

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Gaps

Nugent P;

Baserga R,

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Anti:sense oligo:nucleotide(s) against mutant or native collagen genes for inhibiting collagen expression, e.g for treating osteoarthritis, liver cirrhosis, excessive scarring etc.
                                              Claim 5; Page 24; 55pp; English.
Colige A,
          WPI; 1994-183496/22.
                                                                                                                                                                                                                                                                                                                                                                Guenzburg WH,
                                                                                                                                                                                                                                                                                                WO9709440-A1
                                                                                                                                                                                                                                                                                                                        06-SEP-1996;
                                                                                                                                                                                                                                                                                                                                   06-SEP-1995;
                                                                                                                                                                                                                                     25-MAR-2003
14-NOV-1997
                                                                                                                                                                                                                                                                                                           13-MAR-1997
Prockop D,
                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                          AAT62029;
                                                                                                                                             Query Match
                                                                                                                                                                                                 RESULT 304
                                                                                                                                                                                                        AAT62029
                                                                                                                                                                                                             \texttt{TX} \texttt{EXF} \texttt{EFX} \texttt{EXX} \texttt{X} \texttt{OOOOOOOOOOX} \texttt{Q}
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ö To develop antisense oligos, the test system employed mouse NIH 373 cells stably transfected with an internally deleted construct of the human gene for the pro alpha 1(1) chains of type I procollagen CCLIAI). A series of modified cligos were synthesised using a region at the 3' end of exon I and the first two nucleotides of intron 1 of the exogenous (human) gene as a target. This sequence is given in AAG66555 which corresp. to bys 198 198 225 if the adenthe at the start of transcription is counted as poen. The corresp, sequence of the endogenous (mouse) gene is given in AAG66559. Which corresp to bys 169-195. The antisense oligos are given in AAG66559. Goodely. The antisense oligos inhibit the expression of mutant or normal collagen genes. (Updated on 25-MAR-2003 to correct PN field.) A novel DNA construct (preferably a retroviral vector) has been produced for the treatment of human mammary cell disorders or diseases, including human mammary carcinoma. The DNA construct comprises at least one therapeutic gene under the transcriptional control of the whey acidic Gene expression; human mammary carcinoma cell; whey acidic protein; mouse mammary tumour virus; WAP; MWTV; polymerase chain reaction; ss. expression in human mammary carcinoma cells - using whey acidic 0; Gaps 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; tive 0; Mismatches 3; Indels Murine leukaemia virus retroviral vector BAG PCR primer B. Sequence 20 BP; 1 A; 8 C; 4 G; 7 T; 0 U; 0 Other; BAVA-) BAVARIAN NORDIC RES INST AS. (GSFU-) GSP FORSCHUNGSZENTRUM UMWELT & GESUNDHEI. Saller RM, Salmons B; Example 1; Fig 1; 46pp; English. 26 CGAGGGCTGGGACGAAGA 43 AAT62029 standard; DNA; 20 BP. 95DK-00000976. 96WO-EP003922. 18 cgagggccaagacgaaga (first entry) protein or mouse mammary Local Similarity 83.3 les 15; Conservative (revised) WPI; 1997-192915/17. Gene

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A retroviral vector carrying a DNA sequence encoding SDI-1 (senescent cell derived inhibitor 1), a functional analogue, fragment or antiSense SDI-1 DNA sequence has been developed. The present sequence represents C FOR primer B used in the amplification of mouse leukaemia virus (MLV) retroviral vector beta galactosidase gene (BAG) LTR, for use in the cetroviral vector can be used in the treatment of a polylinker. The retroviral vector can be used in the treatment of disorders or diseases responsive to the anti-proliferative activity of SDI-1, e.g. for the treatment of cencer or restences, septencelly, septencially for the treatment of breast cancer. The retroviral vector acts to introduce the relevant DNA sequences, sense or antisense, into human cells in vitro or in vivo. The retroviral vector antistered by injection or by implantation of a packing cell ine in to the body nearby or at the site of the tumour. (Updated on 25-MAR-2003 to correct PI field.)
                                                                                                                                                                                                                                                                                              ö
protein (WAP) or mouse mammary tumour virus (MMTV) regulatory sequences. The present sequence represents PCR primer B which is involved in the deletion of the U3 region from the murine leukaemia virus (MLV) retroviral vector, known as BAG, and the insertion of a polylinker, which is used in an example for the production of a DNA construct as described above. The WAP and MMTV regularory sequences are able to direct the efficient expression of a linked heterologous gene in primary human mammary cells, including mammary carcinoma cells. (Updated on 25-MAR-2003 to correct PI field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Retroviral vector carrying senescent cell derived inhibitor 1 DNA - used in the treatment of diseases responsive to anti:proliferative activity,
                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MLV; retroviral; vector; senescent cell derived inhibitor 1; SDI-1; antiproliferative; breast cancer; restenosis; human; implantation; tumour; polymerase chain reaction; beta galactosidase gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mouse leukaemia virus retroviral vector BAG gene LTR PCR primer B.
                                                                                                                                                                                                                                                                                              ;
0
                                                                                                                                                                                                                                                   Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.88+02; Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GSF FORSCHUNGSZENTRUM UMWELT & GESUNDHEI.
                                                                                                                                                                                                                 Sequence 20 BP; 7 A; 6 C; 4 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Fig 1; 53pp; English.
                                                                                                                                                                                                                                                                                                                                       64
                                                                                                                                                                                                                                                                                                                                                                                13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP.
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                                                                                                                                                                                                                                                                                                                                         47 CCACCACTCAGAGGAGTC
                                                                                                                                                                                                                                                                                                                                                                                CAATCACTCAGAGGAGAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT85369 standard; DNA; 20
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(first entry)
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11-DEC-1997
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the Arrys758-T92817 represent amplification primers for human genes which are used in the chimeric non-human animal of the invention. The chimeric non-human animal of the invention, preferably a mouse, contains a foreign chromosome (s) or chromosome fragment. The animal is produced by obtaining a hybrid cell by fusion of a cell containing the foreign chromosome with a cell having the ability to form microcells. The microcells are prepared, and fused with cells having differentiative pluripotency and containing the foreign chromosome. These cells are then introduced into an embryo, which is then chromosome. These cells are then introduced into an embryo, which is then chromosome segment to term. The foreign chromosome segment is at least in the long and preferably contains a region for an antibody. The chromosome segment could also contain genes associated with human chromase in the interluckin-2 gene, and the Huntington's disease gene. The expression of foreign genes (especially human genes) in a nonhuman animal is useful for efficient production of proteins, especially of human animal is useful for efficient production of proteins, especially of human animal is useful material can be isolated and fused with myeloma cells to produce the antibody); gene (e.g. to produce the antibody)
                                                                                                                                                                                                                                                                                                                                                                                                                                 PCR primer; amplify; human gene; chimeric non-human animal; antibody; transgenic mouse; chromosome fragment; hybridoma production; microcell; Huntington's disease gene; pluripotent cell; interleukin-2 gene; myeloma cell; immunoglobulin gamma-1; constant region; 1991; ss.
                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                             Primer #2 for immunoglobulin gamma-1 constant region (IGG1).
                                                      Score 13.2; DB 1; Length 20;
Pred. No. 3.8e+02;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hanaoka K, Oshimura M, Ishida I;
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                  Sequence 20 BP; 7 A; 6 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 9; Page 33; 142pp; Japanese.
                                                                                                                                      64
                                                                                                                                                                         2 CAÁTCÁCÍCÁGAGGÁGAC 19
                                                                                                                                                                                                                                                                            AAT92797 standard; DNA; 20 BP.
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96JP-00027940.
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                                                        Query Match
Best Local Similarity 83.3%;
Matches 15; Conservative
                                                                                                                                        47 CCACCACTCAGAGGAGTC
                                                                                                                                                                                                                                                                                                                                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                          05-FEB-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                  AAT92797;
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The present invention describes a method of obtaining pluripotent cells containing foreign chromosomes or their fragments (preferably at least containing foreign chromosomes or their preparing cancerous calls containing the foreign chromosomes or fragments, then fusing these with pluripotent cells such as embryonic stem cells, embryonic reproductive cells embryonic cancer cells or their mutants. Also cascribed are; (1) a method of obtaining hybridoma cells py fusing a cell chart a high ability to produce hybridoma cells (such as mouse A9 cells) with a cell containing the foreign chromosomes or fragments (such as normal human diploid cells); (2) a method of utilising pluripotent cells or produce chimeric and transgenic non-human animals (especially mammals nutroduced; and (3) chimeric animals, their offspring and tissues and cells derived from the offspring produced by a method as in (2). The introduced; and (3) chimeric animals, their offspring and tissues and cells acan be used for the production of monoclonal antibodies for medical use which are of human type and therefore not antigent in humans. They can also be used in the production of chimeric and serve as models for the study of human diseases. Adv52755 to A4v52828 are perve as models for the study of human diseases. Adv52755 to A4v52828 are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ..
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                                                                                                                                                                                                                                                                                                                                                Pluripotent cell; intrinsic gene; chimeric non-human animal; construction; human; antibiotic gene; cancer cell; embryonic; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               non-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pluripotent cells containing foreign chromosomes or fragments - and n
human chimeric animals constructed using them and expressing foreign
genes such as human antibiotic genes.
                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;
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3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hanaoka K, Oshimura M, Ishida I;
83.3%; Pred. No. 3.8e+02;
iive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                               Immunoglobulin gamma-1 constant PCR primer IGG1 #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 4 A; 3 C; 10 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 9; Page 46; 217pp; Japanese.
                                                           364 TCCTCACTTTCCTGGACC 381
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     98WO-JP000860
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          97JP-00062309
                                                                                                 20 récréadegrécade
                                                                                                                                                                                                AAV52794 standard; DNA; 20
      Best Local Similarity 83.3
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tomizuka K, Yoshida H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (KIRI ) KIRIN BEER KK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1998-480821/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9837757-A1
                                                                                                                                                                                                                                                                            27-NOV-1998
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                                                                                                                                                                                                                                     AAV52794;
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3.1%; Score 13.2; DB 1; Length 20;

Query Match

AAZ25788/c

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SOD; superoxide dismutase; antioxidant; superoxide; anion; radical; environmental stress; biological stress; treatment; arthritis; rhemutism; ischaemic heart disease; radiation damage; skin; ageing; cosmetic; plant bioreactor; transgenic plant; expression; herbicide; Basta; resistance; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                            Producing transgenic plants which produce high levels of superoxide
                                                                                                              Basta-resistance (bar) gene PCR primer Bar 2.
                                                                                                                                                                                                                                                                                                        (KOAD ) KOREA ADV INST SCI & TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                Example 4; Page 16; 25pp; English.
                                                                                                                                                                                                                                                                                                                          Kim J, Lee H, Kwon SY, Kwak
                                                                                                                                                                                                                                                                 98KR-00013205.
98KR-00033947.
99KR-00011848.
                                                        BP.
20 TCCTCACCGTCCTGCACC 3
                                                                                                                                                                                                                                                 99EP-00302909.
                                                        AAZ32672 standard; DNA; 20
                                                                                             09-FEB-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Basta-resistant plants
                                                                                                                                                                                                                                                                                                                                            WPI; 1999-582804/50.
                                                                                                                                                                                                                                                                            21-AUG-1998;
06-APR-1999;
                                                                                                                                                                                                                                                 14-APR-1999;
                                                                                                                                                                                                                                                                   14-APR-1998;
                                                                                                                                                                                                           3P952224-A2
                                                                                                                                                                                                                               27-OCT-1999
                                                                                                                                                                                          Synthetic.
                                                                            AAZ32672;
                                       RESULT 308
                                                  AAZ326,
                                                                  셤
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This sequence represents Basta-resistance gene (bar) PCR primer Bar 2, used with primer Bar 1 (AAZ32671) to determine whether the bar gene had been stably introduced into the genome of Basta-resistant plantlets. The pantlets had been previously been transformed with a superoxide dismutase (SOD) expression vector comprising a cucumber fruit dominant promoter (ASO), a cassava mSOD1 gene and the bar gene. SODs are ubiquitous enzymes which convert superoxide anion radicals to hydrogen peroxide is then converted to water by peroxidases or catalases. Superoxide anion radical, are generated by companisms in response to environmental and biological stresses. SOD is thought to be effective in the treatment of arthritis, rheumatism, is chaemic heart disease and radiation damage, and has been used in cosmetics for the prevention of skin ageing The transgenic plants cosmetics for cosmetical for cosmetics, including massage packs, or as an enditive in functional foods or medicines. The plants can be used as
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                                                                          Gaps
                                                                          ;
0
                                  3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; tive 0; Mismatches 3; Indels
Sequence 20 BP; 4 A; 5 C; 7 G; 4 T; 0 U; 0 Other;
                                                                            15; Conservative
                                      Query Match
Best Local Similarity
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214 AGAACTCGGTGGCGGCCA 231
                             8
                             AGATCTCGGTGACGGGCA
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à g RESULT 309

WO9848827-A1

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The present sequence represents a PCR primer for the human p51 gene, which is related to p53 and has cell proliferation regulation and tumour suppression activity. The p51 gene can be used in the investigation, diagnosis and treatment of diseases such as cancer, with which the p53 family cell proliferation regulation is associated. The p51 protein may be used for screening potential agonists and antagonists of its regulatory function, for use as drugs,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ss3 gene; mouse; liver development; signal transduction; liver disease;
tissue repair; cancer; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                        New p53 related human gene p51, useful for diagnosis, investigation and treatment of cancers and screening for potential cell proliferation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                     Human, p51, p53 related gene, cell proliferation, regulation, cancer;
tumour suppression, diagnosis, PCR primer, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 1 A; 4 C; 6 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 3; Page 116; 163pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mouse ss3 gene reverse PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              181 CCAAGGCACATATCCACT 198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20 czaadgecacaaageceaer 3
                                                                                                                                                                                                                                                                                                                                      Obinata M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAV64430 standard; DNA; 20 BP
AAZ25788 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                      98JP-00100467.
                                                                                                                                                                                                                                          99WO-JP001512.
                                                                                                                                                                                                                                                                                          (SAKA ) OTSUKA PHARM CO LTD.
(IKAW/) IKAWA Y.
                                                                             Human p51 PCR primer p51-R6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-MAR-1999 (first entry)
                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                 WPI; 1999-591318/50.
                                                                                                                                                                                                                                                                                                                                      Ikawa Y, Ikawa S,
                                                                                                                                                                                                                                            24-MAR-1999;
                                                                                                                                                                                                                                                                      27-MAR-1998;
                                                                                                                                                Synthetic.
Homo sapiens.
                                                                                                                                                                                       WO9950412-A1
                                                     07-JAN-2000
                                                                                                                                                                                                                  07-OCT-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
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                            AAZ25788;
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Mishra L;

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PCR primers AAZ01426-Z06209 were used to amplify open reading frames (ORFs) of the genome of Chlamydia trachomatis (see AAZ01425). These ORFs encode polypeptides (see AAY36754-4737949) which can be used as vaccines against Chlamydia trachomatis. Antiense and ribozyme sequences can also be used to control growth of the microorganism. Chlamydia trachomatis is responsible for a large number of diseases, e.g. eye diseases such as conventional trachoma, nonendemic trachoma, paratrachoma, and inclusion conjunctivitis; genital diseases such as nongonococcal uretritis, opidymitis, cervicitis, salpingitis, perimpatitis, barbolintis; pneumopathy in breast feeding infants; and veneral lymphogramulomatosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Vaccine; eye disease; conventional trachoma; nonendemic trachoma; paratrachoma; inclusion conjunctivitis; genital disease; perihepatitis; nongonococcal uretritis; epidymitis; cervicitis; salpingitis; PCR primer; bartholinitis; pneumopathy; venereal lymphogranulomatosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR primers AAZ01426-Z06209 were used to amplify open reading frames (ORFS) of the genome of Chlamydia trachomatis (see AAZ01425). These ORFS encode polypeptides (see AAY36794-Y37949) which can be used as vaccines against Chlamydia trachomatis. Antisense and ribozyme sequences can also be used to control growth of the microorganism. Chlamydia trachomatis is responsible for a large number of diseases, e.g. eye diseases such as
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR primer used to amplify an ORF of Chlamydia trachomatis.
                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 4 A; 7 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Genome sequence of Chlamydia trachomatis.
                Genome sequence of Chlamydia trachomatis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 1655; 1755pp; English.
                                                                 Disclosure; Page 1600; 1755pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           351 CTCTACAGCGACTTCCTC 368
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 crcgacaccaarrerre 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         97FR-00015041.
97FR-00016034.
98US-0107077P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ04026 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chlamydia trachomatis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-371125/31.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-NOV-1997;
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04-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ04026;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 312
AAZ04026/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           X#X8X8666666666668
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                  This oligonuclectide is used as a reverse primer, together with a forward primer (see AAV64429), in quantitative FCR of the mouse ss3 gene. 10 demonstrate spairs (see AAV64427-46) were used in quantitative FCR to demonstrate elf (see AAV6411-13), sa3, and 145 (see AAV64414) expression in bloc and liver explant cultures, compared to HNF3-beta, C/EBP, alphafetoprotein and glyceraldehyde 3-phosphate dehydrogenase. The invention provides early developing liver proteins and the genes coding for them (see AAV6410-24). These can be used in the treatment and diagnosis of liver diseases and other disorders, including those relating to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Vaccine, eye disease; conventional trachoma; nonendemic trachoma; paratrachoma; inclusion conjunctivitis; genital disease; perihepatitis; nongonococcal uretritis; epidymitis; cervicitis; salpingitis; PCR primer; bartholinitis; pneumopathy; venereal lymphogranulomatosis; ss.
                                                                                                                                                                                                                                                                    New isolated early liver development genes - used to develop products for treating, e.g. liver disease, hepatocellular carcinoma, degenerative neurological disorders, anaemia, ataxia or haemochromatosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PCR primer used to amplify an ORF of Chlamydia trachomatis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 6 A; 6 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                  Example 2; Fig 20; 92pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    347 GCTGCTCTACAGCGACTT 364
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 97FR-00015041.
97FR-00016034.
98US-0107077P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      oncogenesis and tissue repair
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                                                98WO-US008656.
                                                                                           97US-00841349
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 83.3°
Matches 15, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
Chlamydia trachomatis.
                                                                                                                                                                                                                           WPI; 1999-009382/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (GEST ) GENSET
                                                                                                                                       (MISH/) MISHRA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-NOV-1998;
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04-NOV-1998;
                                                                                           30-APR-1997;
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AAZ03364;

RESULT

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WPI; 1999-371125/31

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Gaps

BP.

8868888888

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Human; glutamic acid decarboxylase; choline acetyltransferase; GAD65; GAD67; ChAT; dopamine receptor; G3PDH; PCR primer; Huntington's disease; neural transplantation; neurological disease; hVT neuron; ss.
                                                                                                Human D2 dopamine receptor PCR forward primer.
AAX00253 standard; DNA; 20
                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                              Freed CR, Kaddis FG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1999-095293/08.
                                                                                                                                                                                                      Synthetic.
Homo sapiens.
                                                                                                                                                                                                                                                       WO9857663-A1
                                                                                                                                                                                                                                                                                                                          17-JUN-1998;
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                                                                   26-MAR-1999
                                                                                                                                                                                                                                                                                         23-DEC-1998.
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                                  AAX00253;
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AAZ98749/c
 셤
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conventional trachoma, nonendemic trachoma, paratrachoma, and inclusion conjunctivitis; genital diseases such as nongonococoal uretritis, epidymitis, cervicitis, salpingitis, perihepatitis, bartholinitis; pneumopathy in breast feeding infants; and venereal lymphogranulomatosis. The polypeptides of the invention may be of use in treating these
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This sequence represents a primer/probe sequence of the invention. The primer and probe sequences are derived from the sequence of the human serine protease gene LKB1, and are used to detect variations in LKB1 leading to Peutz-Jeghers (PD) syndrome. The primers and probes can be used for the diagnosis, investigation and treatment of diseases in which variations in the LKB1 gene are implicated, such as PJ syndrome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LKB1 gene; human; serine protease; Peutz-Jeghers syndrome; PJ syndrome; variation detection; therapy; diagnosis; primer; probe; ss.
                                                                                                                                                                                       Gape
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Primers and probes for use in diagnosis of Peutz-Jeghers syndrome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
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                                                                                                                                                 Match 3.1%; Score 13.2; DB 1; Length 20; Local Similarity 83.3%; Pred. No. 3.8e+02; es 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 3 A; 6 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                    Sequence 20 BP; 1 A; 8 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (CHUG-) CHUGAI RES INST MOLECULAR MEDICINE INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 2; Page 89; 107pp; Japanese
                                                                                                                                                                                                                        225 GCGGCCAAATCGGGAGGC 242
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           89
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                                                                                                                                                                                                                                                        20 GCTGCCAAAGCGGGAGCC 3
                                                                                                                                                                                                                                                                                                                                            AAX79655 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98WO-JP005357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        97JP-00344256
98JP-00280357
                                                                                                                                                                                                                                                                                                                                                                                                                                                Human LKB1 gene primer/probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         51 CACTCAGAGGAGTCTCTG
                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1999-358129/30.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nezu J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                 12-AUG-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      W09928459-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-NOV-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .0-JUN-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Jenne DE,
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Best Local S
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                                                                                       di seases.
                                                                                                                                                      Query Match
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                                                                                                                                                                                         Matches
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RESULT AAX7965

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A method has been developed of treating defective tissue comprising: (i) providing a number of hNT neurons and a neurologically defective mammal having a target tissue comprising defective cells, and (ii) transplanting the hNT neurons into the defective man, so that the neurological defective for the mammal is ameliorated. Also described is a non-human mammal having transplanted hNT neurons. The method is especially used to treat Huntington's disease or other neurological disorders. The method allows the transplanted to of the neurological disorders. The method allows the transplanted to treat and differentiated neurons from cell lines. The present sequence represents a PCR primer used in an example from the present invention for in vitro characterisation of hNT neurons
                                                                                                                                                                                                                                                                                              ,
by
                                                                                                                                                                                                                                                                                    Treatment of neurological disorders, especially Huntington's disease transplantation of differentiated neurons into corpus striatum of affected mammal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; mitochondria; PCR primer; large insert episome; lipofection; epstein barr virus nuclear antigen-1; EBNA-1; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR primer used to amplify human mitochondrial DNA fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 2 A; 10 C; 1 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                        Example 2; Page 36; 53pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   289 AGCTGGTGAAGGACCTGA 306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ВЪ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18 AGATGGTGAAGGACAGGA 1
                                                                                                                   (UYTE-) UNIV TECHNOLOGY CORP.
98WO-US012685.
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RESULT 314 AAX00253/c

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99WO-US019468. 98US-0097961P. 98US-0102691P.

26-AUG-1999;

09-MAR-2000,

WO200012693-A1

UNNC-) UNIV NORTH CAROLINA

01-OCT-1998;

26-AUG-1998;

WPI; 2000-256638/22.

Vos JH;

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Antisense oligonucleotides targeted to nucleic acids encoding human tumor necrosis factor receptor-associated factor (TRAF), useful for treating
                                                                                                                                                                                                                                                                                                                                                                                      The present sequence is that of nested primer NP2, which was used in the amplification of gene fragments obtained from a suppression subtractive hybridization reaction using LAPC xenograft cDNA and designed to identify novel prostate and prostate cancer-specific genes. A 437 bp clone was obtained. Pull-length cDNA (see AAZ94275) was subsequently cloned from a transcription factor that is normally expressed only in testis tissue, but is up-regulated in prostate and other types of cancer. The invention provides diagnostic and therapeutic methods useful in the management of various cancers which express PHELIX, including prostate cancer, bladder cancer, ovarian cancer and testicular cancer
                                                                                                                                                                                                                                                            Testis specific Helix Loop Helix proteins expressed in cancers and useful for the prevention, diagnosis and treatment of prostate, bladder and ovarian tumors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tumour necrosis factor receptor-associated factor; TRAP; human; antisense oligonucleotide; phosphorothioate; antiproliferative; anti-inflammatory; B-selectin; jun kinase; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TRAF2 antisense oligonucleotide ISIS# 16847.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Xu XS;
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                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 31; 62pp; English.
                                                                                                                                                                               Raitano AB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              373 TCCTGGACCGCGACGACG 390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Monia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20 TCCTCGGCGCGGACCACG 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAASSSS6 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99WO-US023171,
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98US-0098610P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cowsert LM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC.
                                                                UROGENESYS INC
                                                                                      AFAR D E.
HUBERT R S.
RAITANO A B.
                                                                                                                                                                               Afar DE, Hubert RS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-303732/26.
                                                                                                                                                                                                                          WPI; 2000-237872/20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    05-OCT-1999;
31-AUG-1998;
31-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Baker BF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAA55556;
                                                                                      (AFAR/)
(HUBE/)
(RAIT/)
                                                                UROG-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This sequence represents a PCR primer used to amplify a fragment of the human mitochondrial DNA. The PCR product is used to create a probe which is used in the Southern blot analysis of cells transfected with the recombinant plasmid of the invention. The plasmid is useful for the production of large insert episomes in mammalian cell. The plasmid comprises a lymphotrophic herpes virus segment and a heterologous insert explication and a hererologous origin of plasmid comprises an episome in both bacterial replication which is maintained as an episome in both bacterial and mammalian cells. The recombinant plasmid is useful for transforming mammalian cells.

Specially B-lymphoblastoid cells (BLC), epithelial cells (BC) or a fusoficient. The recombinant plasmid is also useful for the production of these, by transfecting a mammalian cell with the plasmid by lipofection. The recombinant plasmid is also useful for the production of large-insert episomes in mammalian cells. The invention also relates to an epistein barr virus nuclear antigen-1 (BBNA-1) gene having a partial RS domain deletion which is from 300-700 nucleotides in length. The BBNA-1 gene is useful as a therapeutic agent, especially in gene therapy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PHELIX; human; testis-specific; transcription factor; prostate cancer; bladder cancer; ovary cancer; testicular cancer; gene therapy; diagnosis; vaccine; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                    New recombinant plasmid useful for producing large-insert episomes in mammalian cells comprises a lymphotrophic herpes virus segment linked to a heterologous insert segment.
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0; Gaps

3.1%; Score 13.2; DB 1; Length 20; llarity 83.3%; Pred. No. 3.88+02; Conservative 0; Mismatches 3; Indels

Best Local Similarity Matches 15; Conserv

Query Match

CTGCTCGGTGAAAGCAGA 214

197

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0

CTGCTAGGTGTAAGGAGA

19

BP.

AAZ94278 standard; DNA; 20

RESULT 316

Human PHELIX nested primer NP2

03-JUL-2000 (first entry)

AAZ94278;

99WO-US020137.

31-AUG-1999;

09-MAR-2000

WO200012709-A2

Homo sapiens

Sequence 20 BP; 4 A; 7 C; 3 G; 6 T; 0 U; 0 Other;

Example 2; Page 33; 67pp; English.

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diseases associated with TRAF expression such as inflammatory diseases.

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The present invention relates to antisense oligonucleotides (see AAA55496 -A55757) which are targeted to nucleic acids encoding a human tumour necrobis factor receptor-associated factor (TARF). The antisense enecrobis factor septoral acids encoding a human tumour necrobis factor receptor-associated factor (TARF). The antisense comprise at least one modified internucleotide linkage, which is a phosphorothioate linkage. The oligonucleotides also include at least one modified sugar moiety such as a 2-O-methoxyethy sugar moiety. Sequences AAA55490-A55495 represent nucleotide sequences encoding human TRAF1-6. Included in the invention is a method for treating a human having a disease associated with the expression of TARF comprising administering an antisense oligonucleotide. The reduction of jun kinase administering an antisense oligonucleotide for the reduction of E-coligonucleotide targeted to TRAF-6. A method for the reduction of E-cissues with an antisense oligonucleotide targeted to TRAF-6. The antisense oligonucleotides have antiproliferative and anti-cissues with an antisense oligonucleotide targeted to TRAF-6. The antisense oligonucleotides have antiproliferative and anti-cit in antisense oligonucleotide for the antisense oligonucleotide with an antisense oligonucleotide may also be used as a diagnostic probe for studying gene function
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PTAN; testis specific; prostate cancer; overexpress; chromosome 1q22; diagnose; cancer; breast; vaccine; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                  3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                  Seguence 20 BP; 5 A; 9 C; 5 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR primer (NP2) used in PTAN gene isolation.
                                  Example 16; Page 52; 170pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TGGCCCGCCTGGCGGTGG 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98US-0102556P.
98US-0102910P.
98US-0113229P.
99US-0129518P.
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HUBERT R S.
RAITANO A B.
MITCHELL S C
                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        VO200020589-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-SEP-1999;
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21-DEC-1998;
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(HUBE/)
(RAIT/)
(MITC/)
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PTAN proteins, and sequences encoding them, used for diagnosing and treating cancers, especially breast and prostate cancers.

Mitchell SC

Raitano AB,

Afar DE, Hubert RS,

WPI; 2000-317715/27.

Gaps

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3; Indels

Mismatches

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BPC-1 polypeptides and polymucleotides can be used for the detection of PBC-1 polypeptides and polymucleotides in biological samples, this is particularly useful for detecting cancers expressing BPC-1, e.g. prostate cancer or bladder cancer. Antibodies directed against BPC-1 or antisense This sequence represents a PCR primer used in the isolation of cDNA fragments of the PTAN (testis specific protein expressed in prostate cancer) gene. PTAN is expressed in 3 isoforms PTAN-1, 2, and 3. The PTAN gene is located on chromosome 1022. PTAN is overexpressed in prostate cancer, and has a testis specific expression pattern in adult tissues. PTAN shows no homology to any known gene. PTAN can be used in methods for the diagnosis of cancer, especially prostate or breast cancer, where the normal tissue samples are prostate itssue, or breast tissue, bone tissue, lymphatic tissue, serum, blood, or urine. A vector containing the PTAN rincleotide sequence, a vaccine composition targeting PTAN, PTAN, ribozymes specific for PTAN mRNA and antisense sequences, rospecially breast and prostate cancers. Cancer development can be inhibited by a vaccine composition targeting PTAN New isolated BPC-1 polypeptides, useful for developing products for the diagnosis, staging, prognosis and treatment of cancers, particularly prostate or bladder cancer. BPC-1; oncogene; oncogenic; cancer; prostate; bladder; antibody; antisense; vaccine; detection; prognosis; drug screening; primer; 88. Gaps Primer used for generating human brain specific protein BPC-1 cDNA. ö 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; tive 0; Mismatches 3; Indels Leong K, Raitano AB, Saffran DC; Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other; Example 1, Page 35, 79pp; English. Example 1; Page 31; 71pp; English. 373 TCCTGGACCGCGACGACG 390 m AAZ93048 standard; DNA; 20 BP. 99WO-US018250. 98US-0095982P. 20 recressicación recación (first entry) Conservative UROGENESYS INC Hubert RS, (AFAR/) AFAR D E. (HUBE/) HUBERT R S. (LEON/) LEONG K. (RAIT/) RAITANO A B. (SAFF/) SAFFRAN D C. (JAKO/) JAKOBOVITS A RAITANO A B. SAFFRAN D C. WPI; 2000-206006/18. Local Similarity nes 15; Conserva AFAR D E. HUBERT R S. WO200009691-A2. Jakobovits A; 10-AUG-1999; 10-AUG-1998; 24-JUL-2000 24-FEB-2000. Synthetic. Ë, Query Match UROG-) Best Loca Matches δ g ö

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BP

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22P4F11; human; testis; prostate cancer; diagnosis; gene therapy; marker; vaccine; PCR primer; ss.
                                                                                                                                                                                                                                                                            PCR primer NP2 used in testis-specific 22P4F11 gene amplification.
                                                                                                                                                     AAZ94898 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                      01-AUG-2000
                                                                                                                                                                                              AAZ94898;
                                                                                                                                 AAZ94898/c
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polymuclectides can be used for treating such cancers. The BPC-1 colympetides can also be used in vaccines for treating or inhibiting the development of a cancer expressing BPC-1. The polypeptides and colympetides can also be used for detection, prognosis, drug screening and predicting susceptibility to developing cancer. The BPC-1 polypeptide comprises a CUB domain which is expressed in prostate and bladder cancer carcinoma cells and which shows sequence similarity with CUB domains from other known proteins. In normal human tissues BPC-1 is only expressed in carcinoma cells and which however, it is expressed at high levels in prostate cancer cells and bladder cancer cells. A number of synthetic oligonucleotides were used to generate BPC-1 cDNA from total cell RNA of tumour cells lines. These primers were a cDNA synthesis primer (AAZ93041), two adaptors equences (AAZ93044), AAZ93045), a PCR primer (AAZ93046) and two nested primers (NP)1 used in the amplification method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention relates to a method for obtaining a transgenic plant with male sterility. The method uses site specific recombination to stably transform the plant cells. The method involves the use of DNA encoding the histocyte lethal protein, linked to an anther specific promoter. The method is used to produce male sterile plants. Sequences AAAS9949 to AAAS9961 are used in the method of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          373 TCCTGGACCGCGACGACG 390
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel testes-specific gene 22P4F11 which is expressed in human prostate cancer and is useful as a diagnostic marker and/or therapeutic target for prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cancer, are provided, as well as
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; ative 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                vaccines that prevent development of such cancers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        expressing 22P4F11, especially prostate
                                                                                                                                                                                                                                                                                                                                                                                           Hubert RS, Mitchell SC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Page 28; 54pp; English.
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99US-0146584P.
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Best Local Similarity 83.5%
Conservative
                                                                                                                                                                                                                                                 UROG-) UROGENESYS INC
                                                                                                                                                                                                                                                                         (AFAR/) AFAR D E.
(HUBE/) HUBERT R S.
(MITC/) MITCHELL S C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-303452/26.
WO200018925-A1
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28-JUL-1999;
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ID AAA148
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Gaps

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AAA14807;

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PCR primers AAA14805-07 were used to amplify testis-specific protein Yencoded DNA. The specification describes a new method of diagnosis of prostate cancer. The method comprises determining the level of testis-specific protein Yencoded (TSPY) mRNA or protein, and comparing these presence of elevated TSPY mRNA or protein is indicative of prostate presence of elevated TSPY mRNA or protein is indicative of prostate cancer. Detection of TSPY mRNA expression or protein levels useful in the diagnosis of prostate cancer. Antisense polymucleotides complementary to the coding sequence of human TSPY are useful for treating prostate cancer by inhibiting TSPY transcription (when contacted with the TSPY mRNA). Ribozymes are also useful for treating prostate cancer by cleaving the TSPY mRNA and therefore inhibiting its translation. The vaccine is useful for translation inhibiting the development of prostate cancer in a patient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Foreign chromosome; microcell fusion; homologous recombination; antibody; targeting vector; transgenic animal; disease model; knockout animal; PCR primer; human; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                            Diagnosing prostate cancer by determining the level of testis-specific protein Y-encoded (TSPY) mRNA or protein and comparing these TSPY mRNA or protein levels to those of a normal tissue sample.
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                                                                                          Prostate cancer; testis-specific protein Y-encoded mRNA; TSPY mRNA; vaccine; PCR primer; ss.
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Pred. No. 3.8e+02;
0; Mismatches 3; Indels
                                                               PCR primer for testis-specific protein Y-encoded DNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 20; 32pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             373 TCCTGGACCGCGACG 390
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                                (first entry)
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Best Local Similarity 83.33
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                        (UROG-) UROGENESYS INC
(AFAR/) AFAR D E.
                                                                                                                                                                                                                                                                                                                                                                             Hubert RS;
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AAA09957/c
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Gaps

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The invention relates to a novel method of producing cells containing a modified foreign chromosome or chromosome fragment. The method comprises:

(a) fusing a microcell comprising the foreign chromosome or chromosome fragment, with a cell having a high efficiency for homologous recombination; (b) marking the desired site of insertion of the foreign chromosome using a targeting vector; and (c) inducing deletion or translocation at the marked site. Transpenic animals produced by the method are useful to provide disease models and knockout animals, and in the production of human proteins, particularly human antibodies. This sequence is used in the method of the invention
                                                                                                                                                                                                                                 Producing a cell containing modified foreign chromosomes, useful for the generation of transgenic animals.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            D-pantothenic acid, panB, panC, ilvD, pantotheanate synthatase, ketopantoathydroxymethyltransferase, dihydroxyaciddehydratase; panBC operon, vitamin, primer, ss.
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                                                                                                                                                          Ishida I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 4 A; 3 C; 10 G; 3 T; 0 U; 0 Other;
                                                                                                                                                          Tomizuka K, Yoshida H, Hanaoka K, Oshimura M,
Kuroiwa Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (DEGS ) DEGUSSA-HUELS AG.
(KERJ ) FORSCHUNGSZENTRUM JUELICH GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C. glutamicum panBC operon primer 2.
                                                                                                                                                                                                                                                                              Example 9; Page 68; 316pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sahm H;
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                                                                                                                                                                                                      WPI; 2000-246479/21.
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                                                                                                                              (KIRI ) KIRIN BEER
               WO200010383-A1.
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                                          02-MAR-2000
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Recombinant Corynebacterium DNA useful for production of pantothenic acid vitamin, comprises panB, panC or ilvD genes encoding enzymes.

Example 1; Page 6; 27pp; German.

This invention describes novel recombinant Corynebacterium DNA (I), present in microorganisms of the Corynebacterium genus and comprising at least one of the pans (ketopantohydroxymethyltransferase), panc (pantothenicacidsynthetase), especially the pansC operon, and/or ilvD (dhydroxyaciddshydratase) genes. (I) is useful for the preparation of pantothenic acid a vitamin which has applications including cosmetics, medicine and human and animal nutrition. The new preparation method using fermentation techniques produces the required stereo-isoform D form of pantothenic acid. This sequence represents a primer used in the isolation of the Corynebacterium glutamicum pansC operon which is described in the method of the invention

Sequence 20 BP; 5 A; 8 C; 5 G; 2 T; 0 U; 0 Other;

Query Match
3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels 221 GGTGGCGGCCAAATCGGG 238 N g 8

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Gaps

GTTGTCGGCCACATCGGG 5

RESULT 325 AAA09167/

AAA09167 standard; DNA; 20 AAA09167; 

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Nested primer 2 cloning SSH-generated 36PlA6 gene.

(first entry)

.0-AUG-2000

36PlA6; transcription factor; murine EHF homologue; ETS family; cytostatic; cancer; vaccine; tumorigenesis; primer; ss.

Homo sapiens

WO200020584-A2.

13-APR-2000.

99WO-US022576 12-OCT-1999; 98US-0102744P. 99US-0146447P. 02-OCT-1998; 29-JUL-1999;

UROGENESYS INC. (UROG-) (AFAR/)

(AFAR/) AFAR D E. (HUBE/) HUBERT R S. (MITC/) MITCHELL S C.

Mer DE, Hubert RS, Mitchell SC;

VPI; 2000-303772/26

Novel putative transcription factor gene 36P1A6 for treatment, diagnosis and prevention of prostate, bladder, cervical, ovarian, pancreatic, and colonic cancer.

Example 1; Page 30; 53pp; English.

The human 36P1A6 gene encodes a putative transcription factor based on homology to the murine EHF gene which encodes a transcription factor which is a member of the ETS family. 36P1A6 is expressed in androgendependent and androgen-independent LAPC prostate cancer xenografts and in normal prostate at approximately equal levels. The highest expression is

in the prostate and colon. 36PlA6 may be involved in activating tumorpromoting genes or repressing genes that block tumorigenesis. The 36PlA6 polynucleotides and polympetides are used for the treatment and diagnosis of cancer, e.g. prostate, bladder, cervical, ovarian, pancreatic and colonic cancer (all claimed). Anti-36PlA6 antibodies may be used for purifying 36PlA6 and for isolating 36PlA6 homologues. Antisense purigonal cancer is an electron of the second and transcription and translation of the 36PlA6 gene (claimed). The 36PlA6 polynucleotides and immunogenic fragments may also be used in cancer and polypeptides and immunogenic fragments may also be used in cancer vaccines (claimed)

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Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

ô Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels

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373 TCCTGGACCGCGACG 390 m 20 rccrccccccccace ઠ g

AAC64567 standard; DNA; 20 (first entry) 14-FEB-2001 AAC64567; RESULT 326 AAC64567, 

Human, prostate specific gene, 30P3C8, prostate cancer, diagnosis; cytostatic, gene therapy, vaccine, tumour, primer, 88. Human prostate specific 30P3C8 nested primer 2 SEQ ID NO:25.

Homo sapiens.

WO200061610-A2.

19-OCT-2000.

12-APR-2000; 2000WO-US010218.

12-APR-1999;

(UROG-) UROGENESYS INC.

Raitano AB, Saffran DC Hubert RS, Leong K, Afar DE,

WPI; 2000-619224/59.

30P3CB polypeptide and polynucleotide used for diagnosing, treating and monitoring development of prostate cancer.

Example 1; Page 57; 99pp; English.

The present invention describes human prostate specific protein 30P3C8, which is over-expressed in prostate cancer cells. 30P3C8 has cytostatic content and gene therapy. Methods for activity and can be used in vaccines and gene therapy. Methods for detecting the levels of 30P3C8 protein or mRNA in prostate tissue, bone cativity and can be used in vaccines in an individual or dissegulated cell growth e.g. the presence of cancer in an individual or dissegulated cell growth e.g. hyperplasia. The cancers which are detected or diagnosed are of the bladder, pancreas, colon, brain, bone, lung, kidney or prostate by using cet est samples of serum, blood or urine or tissues of the bladder, pancreas, colon, brain, bone, lung, kidney and prostate approstate cancers lung, winger or prostate by using colon, brain, bone, lung, kidney and prostate. 30P3C8 or polynucleotide sequences can be used for treating cancers expressing colon, brain, bone, lung, mannogenic portions of 30P3C8 are in spaced and other proteins cancers. Immunogenic portions of 30P3C8 are concolonal antibodies bind to 30P3C8 and disrupt interactions between the properties of a receptors for which 30P3C8 is a ligand.

30P3C8 may be a growth factor or other molecule involved in tumour growth and metastasis and other or other molecule involved in tumour growth and metastasis and so anti-30P3C8 antibodies may disrupt the homing or

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invasion or other cancer promoting activities of 30P3C8. The assays are used for detecting, staging and monitoring prostate cancer. The 30P3C8 protein or mRNA are used as additional specific markers for detecting prostate cancer and provide a more specific assay than the serum prostate specific antigen (PSA) assay. The present sequence represents a 30P3C8 nested primer, which is used in the exemplification of the present

G; 2 T; 0 U; 0 Other; Sequence 20 BP; 3 A; 5 C; 10

ö Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels TCTTGGACCGCGACGACG 390 recredecededaceace 3 373 20 à g

Gaps

AAC93282 standard; DNA; 20 BP. AAC93282; RESULT 327 AAC93282 

(first entry) 15-FEB-2001

Human STAT3 phosphorothicate antisense oligonucleotide SEQ ID NO:133

Human; mouse; STAT3; phosphorothioate; antisense oligonucleotide; modulation; signal transducer and activator of transcription; DNA-binding protein; signal transduction; inhibition; apoptosis; inflammatory disease; cancer; antiinflammatory; antirheumatic; cytostatic; immunostimulatory; rheumatoid arthritis; leukaemia; myeloma; melanoma; lymphoma; diagnosis; ss.

Homo sapiens.

WO200061602-A1.

19-OCT-2000.

06-APR-2000; 2000WO-US009054.

99US-00288461. 08-APR-1999;

(ISIS-) ISIS PHARM INC.

Karras JG;

WPI; 2000-619223/59.

New antieense compound for inhibiting the expression of signal transducer and activator of transcription 3 (STAT3) in cells or tissues and treating diseases or condition associated with STAT3, such as rheumatoid arthritis and cancer

Example 12; Page 63; 104pp; English.

The present invention describes an antisense compound (1), 8 to 30 ento-leokages in length, that is targeted to a nucleic acid molecule encoding STAT3 (Signal Transducer and Activator of Transcription) and which inhibits the expression of it. (1) has antiinflammatory, antirheumatic, cytostatic and immunostimulatory activities. (1) is used for inhibiting the expression of STAT3 in cells or tissues, treating an animal having a disease or condition associated with STAT3 or a human having a disease or condition characterised by a reduction in apoptosis, and inducing apoptosis in a cell. Diseases or conditions that are treated are rheumatoid arthritis, cancer of the breast, prostate, brain, had and/or neck, leukaemia, myeloma, melanoma or lymphoma. (I) can also be used for diagnostic methods in detecting and determining the role of STAT3 in various cell functions, physiological processes and conditions and for diagnosting the conditions

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(I) can be used alone or with other drugs as an immunostimulator. (I) is used in sandwich and colourimetric assays, involving enzyme conjugation and radiolabeling and is used in diagnostic kits. AAC93150 encodes human STAT3 and AAC932131 encodes mouse STAT3 as given in the exemplification of the present invention. AAC93151 to AAC93230 and AAC93232 to AAC93239 represent STAT3 phosphorothioate antisense oligonucleotides, and AAC93300 represents a mismatch control oligonucleotide which are used in example
                                                                                                                                                                                                                                                                              Human STAT3 phosphorothioate antisense oligonucleotide SEQ ID NO:134.
                                                                                                                               ..
0
                                                                                                           Score 13.2; DB 1; Length 20; Pred. No. 3.8e+02;
                                                                                                                              3; Indels
                                                                                        Sequence 20 BP; 2 A; 6 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                               0; Mismatches
                                                                                                                                                 136 CCCGCCTGGCGGTGGAGG 153
                                                                                                                                                                     19
                                                                                                                                                                                                                       ВР.
                                                                                                            3.1%;
                                                                                                                                                                CCCGCTTGGTGGTGGACG
                                                                                                                                                                                                                       AAC93283 standard; DNA; 20
                                                                      from the present invention
                                                                                                                                                                                                                                                             (first entry)
                                                                                                                      Best Local Similarity 83.3
Matches 15; Conservative
                                                                                                                                                                                                                                                           15-FEB-2001
                                                                                                                                                                                                                                         AAC93283;
                                                                                                                                                                     N
                                                                                                             Query Match
                                                                                                                                                                                                              AAC93283
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modulation; signal transducer and activator of transcription;
DNA-binding protein; signal transduction; inhibition; apoptosis;
inflammatory disease; cancer; antinflammatory; antirheumatic;
cytostatic; immunostimulatory; rheumatoid arthritis; leukaemia; myeloma;
melanoma; lymphoma; diagnosis; ss. Human; mouse; STAT3; phosphorothioate; antisense oligonucleotide; 06-APR-2000; 2000WO-US009054. #O200061602-A1. Homo sapiens 19-OCT-2000.

08-APR-1999;

(ISIS-) ISIS PHARM INC

Karras JG;

WPI; 2000-619223/59.

New antisense compound for inhibiting the expression of signal transducer and activator of transcription 3 (STAT3) in cells or tissues and treating diseases or condition associated with STAT3, such as rheumatoid arthritis and cancer.

Example 12; Page 63; 104pp; English.

The present invention describes an antisense compound (I), 8 to 30 mucleobases in length, that is targeted to a nucleic acid molecule encoding STAT3 (Signal Transducer and Activator of Transcription) and which inhibits the expression of it. (I) has antiinflammatory, antirheumatic, cytostatic and immunostimulatory activities. (I) is used for inhibiting the expression of STAT3 in cells or tissues, treating an animal having a disease or condition associated with STAT3 or a human having a disease or condition characterised by a reduction in apoptosis, and inducing apoptosis in a cell. Diseases or conditions that are treated are rheumatoid arthritis, cancer of the breast, prostate, brain, head and/or neck, leukaemia, myeloma, melanoma or lymphoma. (I) can also be used for diagnostic methods in detecting and determining the role of STAT3 in various cell functions, physiological processes and conditions

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and for diagnosing the conditions associated with expression of STAT3.

(I) can be used alone or with other drugs as an immunostimulator. (I) is used in sandwich and colourimetric assays, involving enzyme conjugation and radiolabeling and is used in diagnostic kits. AAC93150 encodes human STAT3 and AAC93231 encodes mouse STAT3 as given in the exemplification of represent invention. AAC931310 and AAC93132 to AAC93299 represent statis phosphorothicate antisense oligonucleotides, and AAC93100 represents a mismatch control oligonucleotide which are used in example
                                                                                                                                                                                                                                                              from the present invention
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Seguence 20 BP; 1 A; 7 C; 8 G; 4 T; 0 U; 0 Other;

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Gaps
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Score 13.2; DB 1; Length 20;
Pred. No. 3.8e+02;
0; Mismatches 3; Indels
                       ö
   3.1%;
               Local Similarity 83.3
es 15; Conservative
              Similarity
   Query Match
               Best Loc
Matches
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136 CCCGCCTGGCGGTGGAGG 153 cccecrrecrecrecace 20

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AAC93216 standard; DNA; 20 **AAC93216**/ RESULT

BP

AAC93216;

(first entry) 15-FEB-2001

modulation; signal transducer and activator of transcription;
DNA-binding protein; signal transduction; inhibition; apoptosis;
inflammatory disease; ancer; antiinflammatory; antirheumatic;
cytostatic; immunostimulatory; rheumatoid arthritis; leukaemia; myeloma;
melanoma; lymphoma; diagnosis; ss. Human STAT3 phosphorothicate antisense oligonuclectide SEQ ID NO:67. Human; mouse; STAT3; phosphorothicate; antisense oligonuclectide;

Homo sapiens

#0200061602-A1

19-OCT-2000

06-APR-2000; 2000WO-US009054

99US-00288461. 08-APR-1999;

(ISIS-) ISIS PHARM INC.

Karras JG;

WPI; 2000-619223/59

New antisense compound for inhibiting the expression of signal transducer and activator of transcription 3 (STAT3) in cells or tissues and treating diseases or condition associated with STAT3, such as rheumatoid arthritis cancer and

Example 2; Page 47; 104pp; English

The present invention describes an antisense compound (I), 8 to 30 nucleobases in length, that is targeted to a nucleic acid molecule encoding STAT3 (Signal Transducer and Activator of Transcription) and which inhibits the expression of it. (I) has antiinflammatory, antitheumatic, cytostatic and immunostimulatory activities. (I) is used for inhibiting the expression of STAT3 in cells or tissues, treating an animal having a disease or condition associated with STAT3 or a human having a disease or condition characterised by a reduction in apoptosis, and inducing apoptosis in a cell. Diseases or conditions that are treated are theurist, cancer of the breast, prostate, brain, head and/or neck, lenkaemia, myeloma, melanoma or lymphoma. (I) can also be used for diagnostic methods in detecting and determining the role of 

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and for diagnosing the conditions associated with expression of STRT3.

(I) can be used alone or with other drugs as an imminostimulator. (I) is used in sandwich and colourimetric assays, involving enzyme conjugation and radiolabeling and is used in diagnostic kits. AAC93150 encodes human STAT3 and AAC9321 encodes mouse STAT3 as given in the exemplification of the present invention. AAC93151 to AAC93230 and AAC93232 to AAC93239 represent STAT3 phosphorothinate antisense oligonuclectides, and AAC933300 represents a mismatch control oligonuclectide which are used in example
                                                                                                                                                                                                                                                                from the present invention
                    88999999988888
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Sequence 20 BP; 4 A; 7 C; 3 G; 6 T; 0 U; 0 Other;

ö Score 13.2; DB 1; Length 20; Pred. No. 3.8e+02; 0; Mismatches 3; Indels Query Match
Best Local Similarity 83.3%;
Matches 15; Conservative

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292 TGGTGAAGGACCTGAGCC 309 . H ||||||||||| TGGTGAAGGTGCTGAACC 18 g ઠ

330

BP. AAC93196 standard; DNA; 20

AAC93196;

(first entry) 15-FEB-2001 Human STAT3 phosphorothioate antisense oligonucleotide SEQ ID NO:47.

modulation; signal transducer and activator of transcription; bignal transduction; inhibition; apoptosis; bignal transduction; inhibition; apoptosis; inflammatory disease; cancer; antinflammatory; antirheumatic; cytostatic; immunostimulatory; rheumatoid arthritis; leukaemia; myeloma; melanoma; lymphoma; diagnosis; ss. mouse; STAT3; phosphorothicate; antisense oligonucleotide; 

Homo sapiens

WO200061602-A1.

19-OCT-2000.

06-APR-2000; 2000WO-US009054.

99US-00288461. 08-APR-1999;

PHARM INC.

SISI (-SISI)

Karras JG

WPI; 2000-619223/59.

New antisense compound for inhibiting the expression of signal transducer and activator of transcription 3 (STAT3) in cells or tissues and treating diseases or condition associated with STAT3, such as rheumatoid arthritis and cancer.

Example 2; Page 47; 104pp; English.

The present invention describes an antisense compound (1), 8 to 30 nucleobases in length, that is targeted to a nucleic acid molecule acid molecule acid molecule acid molecule serior system of signal transducer and Activator of Transcription) and which inhibits the expression of it. (1) has antiinflammatory, antirheumatic, cytostatic and immunostimulatory activities. (1) is used for inhibiting the expression of STRT3 in cells or tissues, treating an animal having a disease or condition associated with STRT3 or a human having a disease or condition characterised by a reduction in apoptosis, and inducing apoptosis in a cell. Diseases or conditions that are treated are theumatoid arthritis, cancer of the breast, prostate, brain, head and/or neck, leukaemia, myeloma, melanoma or lymphoma. (1) can also be

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used for diagnostic methods in detecting and determining the role of STAT3 in various cell functions, physiological processes and conditions and for diagnosing the conditions associated with expression of STAT3.

(I) can be used alone or with other drugs as an immunostimulator. (I) is used in sandwich and colourimetric assays, involving enzyme conjugation and radiolabeling and is used in diagnostic kits. AAC93150 encodes human STAT3 and AAC93221 encodes mouse STAT3 as given in the exemplification of the present invention. AAC93151 to AAC93230 and AAC93232 to AAC93239 represent STAT3 phosphorothicate antisense oligonucleotides, and AAC93300 represents a mismatch control oligonucleotide which are used in example from the present invention

Sequence 20 BP; 2 A; 5 C; 9 G; 4 T; 0 U; 0 Other;

Gaps ö Score 13.2; DB 1; Length 20; Pred. No. 3.8e+02; 0; Mismatches 3; Indels ö 3.1%; Conservative Similarity Local S... 15; Query Match Matches

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ccceccreecegreeAge 153 1 cccccrrccrccrccrccrcc 18 136 g ઠે

RESULT

AAC64486 standard; DNA; 20 BP

Prostate tumour associated gene 24P4C12 nested primer 2 SEQ ID NO:41.

(first entry)

13-FEB-2001

AAC64486;

Human; prostate tumour associated gene; 24P4Cl2; prostate cancer; transmembrane protein; diagnosis; anticancer; cytostatic; vaccine; gene therapy; PCR primer; ss.

Homo sapiens

WO200061746-A1.

19-OCT-2000.

12-APR-2000; 2000WO-US010039.

99US-0128858P. 12-APR-1999;

(UROG-) UROGENESYS INC

Saffran DC; Raitano AB, Leong K, Afar DE, Hubert RS,

WPI; 2000-672681/65.

Novel 24P4C12 polypeptides and polynucleotides, used in the diagnosis and treatment of cancer, especially prostate cancer.

Example 1; Page 65; 137pp; English.

The present invention describes a prostate tumour associated gene, designated 24P4C12, and its encoded protein. 24P4C12 has anticancer and cytostatic activity, and can be used in vaccine production and in gene therapy. A pharmaceutical composition or vaccine comprising 24P4C12 can be used to treat a patient with cancer, especially prostate cancer, the vaccine can also be used to inhibit the development or progression of cancer. The polypeptides and polymucleotides can be used to diagnose cancers, especially prostate cancer. A transgenic animal comprising 24P4C12 can be used for the development and screening of therapeutic reagents. The polypeptide is a transmembrane protein which is expressed specifically in prostate cancer, allowing the development of more specific anticancer therapies and diagnostic assays

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention provides the protein and coding sequences of human cancer related protein 20P2H8. The gene, which is found at chromosome 15q32-23, is upregulated in cancers such as that of the prostate, bladder, colon and pancreas. The sequences can be used to diagnose and treat these cancers, and to vaccinate against them. The present sequence is a PCR primer for the coding sequence of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                         Gaps
                                                                                                                                                                                                                                         Human, cancer related protein 20P2H8; vaccine, chromosome 15q32-23; prostate cancer; bladder cancer; colon cancer; pancreatic cancer; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20P2H8 polynucleotides and polypeptides useful for diagnosing and treating cancer, and for screening for screening for modulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Jakobovits A;
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Length 20;
                         Indels
                                                                                                                                                                                                                 Human cancer related protein 20P2H8 cDNA PCR primer #3
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Score 13.2; DB 1;
Pred. No. 3.8e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene fragment amplifying NP2 primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TCCTGGACCGCGACGACG 390
                                                   373 TCCTGGACCGCGACGACG 390
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                            15; Conservative
                                                                                                                                          AAF85709 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Raitano AB,
                                                                                                                                                                                                                                                                                                                                                                                                                          (UROG-) UROGENESYS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-308645/32.
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              Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                  28-OCT-1999;
                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                           10-DEC-2001
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                                                                                                                                                                                                                                                                                                                                                 03-MAY-2001
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                                                                                                                                                                   AAF85709;
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Best Local &
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    Query Match
                            Matches
                                                                                                                   RESULT 332
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The present invention relates to methods and compositions for the diagnosis and therapy of prostate cancer which utilise human SGP28 (Specific granule protein 28) gene and proteins. The method involves detecting cancers, particularly of prostate and colon, from expression of SGP28 protein. The expression of SGP28, which is an extractly up-regulated in prostate tumours. SGP28 sequence is used for diagnosis (including in vivo imaging), staging, monitoring and prognosis of prostatic and colon cancer, and for assisting selection of therapy.

Composition or vaccine that contains a vector expressing an antibody specific for SGP28 expressing cancers can be treated by administering a composition or vaccine that contains a vector expressing an antibody specific for SGP28 protein or therapeutic agent. SGP28 grotein or its gracial conjugated to toxin or therapeutic agent. SGP28 grotein or its specific for SGP28 protein or therapeutic agent. SGP28 grotein or its also used as source of therapeutic antisense or ribozyme agents. SGP28 protein and its creaments and for isolating related sequences. SGP28 protein and its conjudence and solation and its calcium entry into prostatic cells, for recombinant production of SGP28 peptides and for isolating related sequences. SGP28 protein and its creaments are used to raise specific antibodies (Ab) and to identify specific binding agents (potentially useful as therapeutic and diagnostic agents) and also potential anticancer agents. The present sequence is a nested primer 2 (NP2) used to amplify gene fragments resulting from SSH (suppression subtractive hybridisation) reaction. This sequence is used in the SSH isolation of cDNA fragment of human SGP28 gene
Human, specific granule protein 28; SGP28; therapy; PCR primer; prostate; colon; cancer; prognosis; vaccine; anticancer; SSH; suppression subtractive hybridisation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Detecting cancers, particularly of prostate and colon, from overexpression of SGP28 protein, also methods for treating these cancers e.g. by vaccination with the protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                             Afar DEH, Mitchell SC, Faris M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               49.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human AKAP10 coding sequence PCR primer SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Seguence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 59; 102pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  373 TCCTGGACCGCGACGACG 390
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                                                                                                                                                                                                                  27-OCT-2000; 2000WO-US029607
                                                                                                                                                                                                                                                              99US-0162610P
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                                                                                                                                                                                                                                                                                                                                               Raitano AB,
                                                                                                                                                                                                                                                                                                    (UROG-) UROGENESYS INC
                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-308685/32.
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                                                                                                                               WO200131343-A2.
                                                                                                                                                                                                                                                                                                                                               Hubert RS, R
Jakobovits A;
                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                           28-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12-JUN-2001
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Matches
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Producing a database for identifying polymorphic genetic markers, comprises obtaining data relating to members of a healthy population and entering the information into a database.
                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention provides a database of human samples obtained from healthy individuals which can be used to identify polymorphic genetic markers. Data obtained for the database can be used to sort the samples by parameters such as age, sex and ethnicity. This is useful in linking markers with diseases, susceptibility to infection and drug responses. The present primer was used in an assay to demonstrate the uses of the database of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human T-type low voltage activated calcium channel alphalG-c; stress; epilepsy; schizophrenia; depression; sleep disorder; Cushing's disease; endocrine disorder; respiratory disorder; peripheral muscle disorder; muscle excitability; fertilisation; contraception; hypertension; neuronal firing regulation; cardiovascular disorder; gene therapy; forensic analysis; epidemiological study; neuroleptic; PCR primer; ss.
SNP; human; genetic marker; disease; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   / Match 3.1%; Score 13.2; DB 1; Length 20; Local Similarity 83.3%; Pred. No. 3.8e+02; nes 15; Conservative 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                             Rodi
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                                                                                                                                                                                                                                                                             Koester H, Van Den Boom D,
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Database; polymorphism; SNP; h
drug response; PCR primer; ss.
                                                                                                                                                                  13-OCT-1999; 99US-0159176P.
10-UUL-2000; 2000US-0217251P.
10-UUL-2000; 2000US-0217658P.
19-SEP-2000; 2000US-00663968.
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                                                                                                                                      13-OCT-2000; 2000WO-US028413
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                               (SEQU-) SEQUENOM INC
                                                                                                                                                                                                                                                                                                                         WPI; 2001-273865/28.
                                                                                                                                                                                                                                                                                             Jurinke C;
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                                                                             WO200127857-A2
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                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-MAY-2001.
                                                                                                          19-APR-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAD04754;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 m
                                                                                                                                                                                                                                                                               Braun A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                               Chiu N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 335
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Matches
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Gaps

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WPI; 2001-308646/32

Erlander MG;

Zhu JY,

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The invention relates to isoform of human T-type low voltage activated calcium channel (alpha1G-c) cDNA and protein. Cells transformed with calcium channel DNA to express calcium alpha1G-c channel protein are used calcium channel DNA to express calcium alpha1G-c channel protein are used to identify specific modulators (antagonists or agonists). These condulators are useful as therapeutic agents and are used for treating wide range of calcium alpha1G-c channel-mediated disorders, e.g. stress conductive disorders, respiratory disorder, peripheral muscle disorder, endocrine disorders, respiratory disorder, peripheral muscle disorder, contraception, disorders involving the muscle excitability, fertilisation, contraception, disorders involving cipalals and cardiovascular disorders (e.g. atherosclerosis, cardiac hypertrophy, angina pectoris). Calcium alpha1G-c channel DNA is useful contionally used as antisense sequences, in gene therapy. Calcium channel contionally used as antisense sequences, in gene therapy. Calcium channel contionally used as antisense sequences, in gene therapy. Calcium channel contionally and identifying related muschindard hybridistation or immunological assays. The present sequence is 18341P PCR primer used for immunological assays. The present sequence is 18341P PCR primer used for thalamus library for generating human calcium alpha1G-c channel sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; 36P6D5 protein; secreted tumour antigen; therapy; cancer; kidney; bladder; ovary; breast; pancreas; colon; lung; vaccine; cytostatic; SSH; suppression subtractive hybridisation; PCR primer; ss.
                                                                                                                                                New nucleic acid encoding human calcium channel protein, useful for identifying specific modulators and potential pharmaceuticals for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human 36P6D5 gene fragment amplifying primer NP2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 3 A; 6 C; 9 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3.1%; Score 13.2; DB 1;
Local Similarity 83.3%; Pred. No. 3.8e+02;
les 15; Conservative 0; Mismatches 3;
                                                                                                                                                                                                                                 Example 2; Page 45; 115pp; English.
                                                                          Pyati J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3 ACTGCCAGTGGCCGAGGG 20
                                       (ORTH ) ORTHO-MCNEIL PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAD04811 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-OCT-2000; 2000WO-US029894.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99US-0162417P.
  99US-00426998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14 ACTGCGGGTGACCGAGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Jakobovite A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                            treating e.g. epilepsy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (UROG-) UROGENESYS INC.
                                                                          Galindo JE,
                                                                                                               WPI; 2001-300486/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200131015-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        03-MAY-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Raitano AB,
Witchell SC;
  26-OCT-1999;
                                                                            Dubin AE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAD04811;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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Matches
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The present invention relates to a gene and its encoded secreted tumour antigen, termed 36P6D5. These sequences are used for the diagnosis and treatment of various cancers which express 36P6D5, such as cancers of the kidney, bladder, ovary, breast, pancreas, colon and lungs. In normal individuals 36P6D5 protein, is predominantly expressed in pancreas, with comprising immunogenic protein of 36P6D5 is useful for inhibiting the comprising 36P6D5 protein is useful for diagnosis and/or prognosis of prostate cancer and other cancers, for modulating or inhibiting the comprising 36P6D5 genes and/or translation of the 36P6D5 transcripts, and as therapeutic agents. The present sequence is a nested primer (NP)2 used to amplify gene fragments resulting from SSH (suppression to counter bybridisation) traction. This sequence is used in the SSH isolation of cDNA fragment of human 36P6D5 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; PC-LECTIN; C-type lectin; transmembrane antigen; normal testis;
layllin homologue; prostate cancer antigen; overexpression;
androgen-dependent prostate cancer; diagnosis; prognosis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New PC-LECTIN polynuclectide encoding a transmembrane antigen over expressed in human prostate cancer, useful for the prognosis, diagnosis and treatment of prostate cancer.
                                           Detecting presence of cancer expressing 36P6D5 protein in individual by comparing protein level in test sample to normal sample, where elevated level of protein in test sample indicates presence of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primer NP2, SEQ ID NO:18, used in human PC-LECTIN cDNA isolation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a novel human C-type lectin transmembrane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 3 A; S C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Raitano AB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Jakobovits A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Page 59; 116pp; English.
                                                                                                                    Example 1; Page 70; 113pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             373 TCCTGGACCGCGACG 390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20 TCCTCGCCCCGACCACG 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          012/c
AAF76012 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (UROG-) UROGENESYS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Afar DEH, Hubert RS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-211222/21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
hes 15; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200112811-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-FEB-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22-MAY-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAF76012;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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Matches
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Gaps

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Length 20; 3; Indels Hubert RS;

Afar DEH,

Faris M,

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contigen, PC-LECTIN (AAB71309) and cDNA encoding it (AAF76004). The expression of the human PC-LECTIN gene is normally restricted to the expression of the human PC-LECTIN gene is normally restricted to the captures and indicate the cancer. PC-LECTIN therefore androgen-independent prostate tumours, and expression is therefore likely cobe dependent prostate tumours compared with androgen-independent prostate cancer. Human PC-LECTIN therefore represents a diagnostic and therapeutic target for prostate cancer. Coparticularly androgen-dependent prostate cancer. Human PC-LECTIN exhibits compared to be the human orthologue of laylin, as diverges significantly in a key functional domain proposed for the laylin coppered to protein. Human PC-LECTIN or nimmunogenic portion thereof, a vector coconic thuman PC-LECTIN antisense nucleotide, a PC-LECTIN concleotide-targetted ribozyme, or an anti- PC-LECTIN antibody may be used to protein-targetted ribozyme, or an anti- PC-LECTIN antibody may be used to protein-targetted ribozyme, or an anti- PC-LECTIN antibody may be used to prepare a composition for treating a patient with a cancer. Colon, pancreatic, testicular, carvical or ovarian cancers that express colon, pancreatic, testicular, carvical or ovarian cancers that express pC-LECTIN proteins are also useful for diagnosing the presence colon, pancreatic, testicular, carvical or ovarian cancers that expressing cancers. PC-LECTIN proteins are also useful for prograte cancer and other PC-LECTIN-expressing cancers. PC-LECTIN corporate cancer and other PC-LECTIN-expressing cancers. PC-LECTIN cancers are useful in the concept of dentify molecules may additionally be used in drug discovery to dentify molecules that modulate PC-LECTIN function or expression. The present sequence represents a PCR primer used in the isolation of human processing cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PC-LECTIN CDNA
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8888888888888888888888888888888

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

Gaps ö Score 13.2; DB 1; Length 20; Pred. No. 3.8e+02; 0; Mismatches 3; Indels ch 3.1%; l Similarity 83.3%; 15; Conservative Query Match Best Local Similarity Best Loc Matches

373 TCCTGGACCGCGACGACG 390 m Tocroscoccoacaca 20 a ð

AAD01985 standard; DNA; 20 RESULT 338 AAD01985/c

BP

26-MAR-2001 AAD01985; 

(first entry)

TCV 12 oligonucleotide to construct pMOG845 plasmid.

TPS; TPP; bipartite enzyme; trehalose phosphate synthase; trehalose; trehalose phosphatase; transgenic plant; stress resistance; cold; drought; natural flavour; stabiliser; forced water extraction; freeze drying; nutritional value; ss.

Jnidentified.

AU200048921-A

26-OCT-2000

31-JUL-2000; 2000AU-00048921.

97AU-00010085. 09-JAN-1997;

(MOGE-) MOGEN INT NV.

Krutwagen RWHH, Voogd E; Goddijn OJM, Verwoerd TC, WPI; 2001-007580/02. Chimeric gene encoding bipartite trehalose synthesis enzyme, useful for producing transgenic plants with increased trehalose content.

Disclosure; Page 16; 59pp; English

The present invention relates to a chimeric gene comprising a potato
patatin promoter and proteinase inhibitor II terminator [PotPiII],
cencoling bipartite trehalose synthesising enzyme and method for
production of trehalose and increasing the level of trehalose
production of trehalose. This bipartite enzyme with trehalose
centendes by trehalase. This bipartite enzyme with trehalose phosphate
cynthase (TPS) and trehalose phosphate phosphates (TPP) activities,
chances the production of trehalose as it enables the completion of
centended to a trehalose and glucose-6-phosphate into trehalose
cat one and the same site. Plants that contain chimeric gene have improved
cat one and the same site. Plants that contain chimeric gene have improved
cat one and the same site. Plants that contain chimeric gene have improved
cat one and the same site. Plants that contain chimeric gene have improved
cresistance to stress (cold or drought) and better post-harvest quality
contains shelf-life. Trehalose is used for forced water extraction, of
creention of natural flavours and mutritional value and allowing rapid
creconstitution, also as e.g. a stabiliset for vaccines, enzymes,
constitution, also as e.g. a stabiliset for vaccines, enzymes,
constitution, also as e.g. a stabiliset for vaccines, chemically inert
contains the present sequence is a TCV 12 oligonucleotide used in the construction of pMOG845 plasmid

Seguence 20 BP; 4 A; 6 C; 5 G; 5 T; 0 U; 0 Other;

Gaps ö 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; tive 0; Mismatches 3; Indels 15; Conservative Similarity Query Match Local Matches

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RESULT 339 AAF83890/

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890/c AAF83890 standard; DNA; 20 AAF83890;

BP

(first entry) 06-AUG-2001 Nested primer (NP)2 used in human PHOR-1 cDNA isolation.

G-protein-coupled receptor; prostate; cancer; PHOR-1; kidney; uterine; cervical; stomach; rectal; cytostatic; vaccine; cell function regulator; human; prostate homologue of olfactory receptor-1; PCR primer; ss.

Homo sapiens.

WO200125434-A1 12-APR-2001 

05-OCT-2000; 2000WO-US027543

99US-0157902P. 05-OCT-1999;

(UROG-) UROGENESYS INC

Jakobovits A, Faris M, Hubert Raitano AB, Afar DEH, J Mitchell SC, Saffran DC;

RS;

WPI; 2001-367230/38.

Novel gene designated PHOR-1, a G-protein-coupled receptor up-regulated in prostate cancer, useful as diagnostic marker and therapeutic target for cancers of prostate, kidney, uterus.

Example 1, Page 59, 139pp; English.

The invention relates to a novel G-protein-coupled receptor up-regulated in prostate cancer, termed PHOR-1. The encoding cDNA is contained in plasmid designated pl01P3A11 deposited with ATCC as Accession No.PTA-312. PHOR-1 polypeptides and polymucleotides are useful for diagnosing the presence of cancer, especially prostate, kidney, uterine, cervical,

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25-SEP-2001
                                                                                                                                                                                                                                                                                                                                      AAD12168;
                                                                                                                                                                                                                                                                                                    RESULT 341
                                                                                                                                                                                                                                                                                                             AAD12168,
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stomach or rectal cancer by determining and comparing the level of the protein or mRNA expression in teet and normal tissue samples. Pharmaceutical compositions comprising PHOR-1 is useful for treating cancer. PHOR-1 proteins are useful for identifying ligands and other agents and cellular constituents that binds to PHOR-1 gene product and for generating antibodies which are useful in diagnostic, prognostic and imaging methodologies and for the treatment of prostate cancer. Cell inse expressing PHOR-1 are useful for identifying protein-protein interactions mediated by PHOR-1. The present sequence represents a primer used in isolation of the PHOR-1 (prostate homologue of olfactory receptor
                                                                                                                                                                                                                                                                                                   Human, PTP1B; protein phosphatase 1B inhibitor; antisense; gene therapy; infection; inflammation; tumour; prophylaxis; phosphorothioate; ss.
                                                                                                                                               Gaps
                                                                                                                                              <u>;</u>
                                                                                                                             Length 20;
                                                                                                                                              3; Indels
                                                                                                                                                                                                                                                                                   Human PTP1B antisense oligonucleotide (ISIS# 107769).
                                                                                                           Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                           note = "Phosphorothioate backbone"
                                                                                                                          Score 13.2; DB 1;
Pred. No. 3.8e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                     1. .5
/*tag= b
/mod base= OTHER
/note= "Methoxyethyl residues"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             note= "Methoxyethyl residues"
8. .20
                                                                                                                                                                                                                                                                                                                                                         ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                           /*tag= a
'mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /*tag= c
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6. .9
/*tag= e
/mod_base= m5c
14. .16
/*tag= f
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /mod_base= m5c
16. .20
/*tag= c
                                                                                                                                                                373 TCCTGGACCGCGACG 390
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'mod_base= m5c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /*tag= g
/mod_base= m5c
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                                                                                                                                                                                                                             AAD11960 standard; DNA; 20 BP.
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                                                                                                                           Ouery Match
Best Local Similarity 83.3%;
Matches 15; Conservative
                                                                                                                                                                           20 TCCTCGGCGGCGACCACG
                                                                                                                                                                                                                                                                 (first entry)
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/*tag=
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                                                                                                                                                                                                                                                                                                                               Homo sapiens.
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                                                                                                                                                                                                                                                                 25-SEP-2001
                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                               AAD11960;
                                                                                          -1) CDNA
                                                                                                                                                                                                            340
                                                                                                                                                                                                           RESULT 34
AAD11960
 88888888888888888
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                                                                                                                                                                                                                                                                                                   The invention is directed to antisense compounds, particularly oligonucleotides which are targetted to a DNA encoding protein phosphatase 18 (PPTB) to modulate its expression. The antisense compounds are useful for diagnosis, prophylaxis and treatment of diseases associated with the expression of PTP1B, to prevent or delay infection, inflammation and tumour formation and as a research reagent. The PTP1B DNA is useful in gene therapy. The present sequence is an antisense oligonucleotide with a phosphorchioate backbone. This oligo is targetted to human PTP1B to inhibit its expression
                                                                                                                                                New antisense compounds capable of modulating expression of human protein phosphatase 1B, useful for diagnosis, prophylaxis and treatment of diseases associated with expression of protein phosphatase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rat; PTP1B; protein phosphatase 1B inhibitor; antisense; gene therapy; infection; inflammation; tumour; prophylaxis; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rat PTP1B antisense oligonucleotide (ISIS# 111615).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 2 A; 11 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mod_base= OTHER
/note= "Phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'note = "Methoxyethyl residues"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /*tag= g
/mod_base= m5c
16..20
/*tag= c
/mod_base= OTHER
                                                                                                                                                                                                                                                  Example 15; Col 42; 71pp; English.
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/*tag= b
/mod_base= OTHER
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/mod_base= m5c
10. .11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  303 CTGAGCCCCGGGGACCGC 320
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'mod_base= m5c
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(ISIS-) ISIS PHARM INC.
                                                   Cowsert LM, Wyatt J;
                                                                                                 WPI; 2001-432181/46.
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Synthetic.
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The invention is directed to antisense compounds, particularly oligonuclectides which are targetted to a DNA encoding protein phosphatase 1B (PPIB) to modulate its expression. The antisense expension of modulate its expression. The antisense associated with the expression of PTPIB, to prevent or delay infection, inflammation and tumour formation and as a research reagent. The PTPIB DNA is useful in gene therapy. The present sequence is an antisense oligonuclectide with a phosphorchicate backbone. This oligo is targetted to rat PTPIB to inhibit its expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Zebrafish; morphogenic signal; neuron; hedgehog gene; embryonic patterning; cell culture; cell differentiation; ischaemia; cell proliferative disorder; intracerebral grafting; Huntington's chorea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New antisense compounds capable of modulating expression of huma
phosphatase 1B, useful for diagnosis, prophylaxis and treatment
diseases associated with expression of protein phosphatase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Zebrafish hedgehog gene amplifying degenerate PCR primer, hh5.2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.88+02; tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 2 A; 6 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                       note= "Methoxyethyl residues"
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/*tag= g
/mod_base= m5c
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                                     '*tag≈ e
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/mod_base=
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Best Local Similarity 83.3
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention is directed to antisense compounds, particularly oligonucleotides which are targetted to a DNA encoding protein phosphatase 1B (PPTB) to modulate its expression. The antisense encompounds are useful for diagnosis, prophylaxis and treatment of diseases associated with the expression of PTP1B, to prevent or delay infection, inflammation and tumour formation and as a research reagent. The PTP1B DNA is useful in gene therapy. The present sequence is an antisense oligonucleotide with a phosphorothicate backbone. This oligo is targetted to rat PTP1B to inhibit its expression
                                                                                                                                                                                                                                                                                                                                                 New antisense compounds capable of modulating expression of human protein phosphatase 1B, useful for diagnosis, prophylaxis and treatment of diseases associated with expression of protein phosphatase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rat, PTP1B; protein phosphatase 1B inhibitor; antisense; gene therapy; infection; inflammation; tumour; prophylaxis; phosphorothioate; ss.
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**tag= a
**dase= OTHER
'note= "Phosphorothicate backbone"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'note= "Methoxyethyl residues"
/note= "Methoxyethyl residues"
16. .1,
*tag= h
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Best Local Matches 1

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RESULT 342 AAD121

protein

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Gaps

Sequence 20 BP; 3 A; 6 C; 2 G; 1 T; 0 U; 8 Other; on 11-SEP-2003 to standardise OS field) AAC887 ద g ઠ The present invention relates to assay for screening compounds that potentiate or inhibit binding of hedgehog polypeptide to naturally occurring patched receptor. The hedgehog protechs comprise morphogenic signals produced by embryonic patterning centres, and are involved in the formation and maintenance of ordered spatial arrangements of formation and maintenance of ordered spatial arrangements of different discussions in the proteins can be used to generate and/or maintain an array of different vertebrate tissues both in vitro and in vivo. The invention also relates to a method for modulating growth, differentiation or survival of a mammalian cell (e.g. neuron, testicular cell) responsive to hedgehog agonists and antegonists can be used in cell culture techniques to enhance survival and maintenance of neurons and various vertebrate organogenic pathways. The hedgehog gene is useful in determining whether a patient is at the risk of disorder characterised by commend cell proliferation or aberrant control of differentiation. The Screening compounds that potentiate or inhibit binding of hedgehog polypeptide to naturally occurring patched receptor, comprises contacting polypeptide with receptor and test compound, and detecting change in binding. hedgehog proteins or mimetics can be used to induce foetal neurons especially neuronal stem cells in intracerebral grafting. The protein or its mimetic can be used in the treatment of neurological conditions e.g. injury to nervous system, ischaemia resulting from stroke, Alzheimer's disease, Parkinson's disease, Huntington's chorea, amyotrophic lateral sclerosis (ALS) and multiple sclerosis. The present sequence is a degenerate PCR primer used to amplify Zebrafish hedgehog gene. (Updated neurological disorder, Alzheimer's disease; Parkinson's disease; amyotrophic lateral sclerosis; ALS; multiple sclerosis; PCR primer; ss. Mcmahon AP (IMCR ) IMPERIAL CANCER RES TECHNOLOGY LTD. (HARD ) HARVARD COLLEGE. Location/Qualifiers Example 4; Col 91; 127pp; English. Ingham PW, 93US-00176427. 94US-00356060. 95US-00435093. 95US-00460900. 95US-00462386. 96US-00674509. /*tag= f /mod_base= i mod_base= i mod_base= mod_base= mod_base= mod_base= Ω υ Ъ ø tag= *tag= *tag= *tag= *tag= Tabin CJ, WPI; 2001-440859/47 Key modified_base 30-DEC-1993; 14-DEC-1994; 04-MAY-1995; 05-JUN-1995; modified_base modified_base modified base modified base modified base JS6261786-B1 02-JUL-1996; Danio rerio 17-JUL-2001 Marigo V,

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nucleic acid or its fragments, useful for diagnosing and treating cancer and neurological disorders, corresponds to a catenin-binding protein in signal transduction and gene regulatory pathways.
                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Human catenin-binding zinc finger protein PCR primer FVR359R.
Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 60.0%; Pred. No. 3.8e+02; Matches 12; Conservative 2; Mismatches 6; Indels
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                                                                                                     133 TGGCCCGCCTGGCGGTGGAG 152
                                                                                                                                                  20 TNGCNMGNYTNGCNGTNGAG 1
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Best Local Similarity 83.35,
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Length 20; 6; Indels

3.1%; Score 13.2; DB 1; 60.0%; Pred. No. 3.8e+02; tive 2; Mismatches 6;

Query Match 3.1' Best Local Similarity 60.0 Matches 12; Conservative

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Sequence 20 BP; 3 A; 6 C; 2 G; 1 T; 0 U; 8 Other;
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The invention relates to nucleic acids encoding hedgehog proteins selected from sonic hedgehog (Shh), indian hedgehog (Ihh), desert hedgehog (Dhh) polypepitales. The hedgehog genes are involved in the formation of ordered spatial arrangements of differentiated tissue in vertebrates. The nucleic acid sequences are useful for producing hedgehog proteins, used for promoting differentiation of, or survival of differentiation on unronal cells, and for promoting proliferation, survival or differentiation of mesenchymal, endodermal or ectodermal tissue, particularly chondrocytes, or testicular germ line cells. Sequences AAH76125-126 represent PCR primers for amplifying a zebrafish Shh genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel nucleic acid encoding a hedgehog polypeptide, used to produce the polypeptide, which is used to promote proliferation, survival, and/or differentiation of neuronal and mesodermal tissue.
                                                 Hedgehog protein; sonic hedgehog; Shh; indian hedgehog; Ihh; Dhh; desert hedgehog; cell differentiation; zebrafish; PCR primer; ss.
Zebrafish Shh DNA amplifying primer hh 3.3.
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(IMCR ) IMPERIAL CANCER RES TECHNOLOGY LTD.
                                                                                                                                                                                                           Location/Qualifiers
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/mod_base= i
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95US-00435093.
95US-00462386.
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                                                                                                                                                                                                         Key
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14-DEC-1994;
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                                                                                                                                                          Danio rerio.
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                                                                                                                              Synthetic
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tumour; kidney; brain; bone; ovary; breast; pancreas; uterus; colon; lung; cyrostatic; gene therapy; ancibody therapy; ribozyme; liver; single chain monoclonal antibody; serum; blood; urine; bladder; cervix; rectum; stomach, human; chromosome 1q31-q32. 83P5G4; PCR primer; DNA adaptor; prostate; testis; tissue; cancer; ss;

Human prostate-related gene 83P5G4 cDNA nested primer #2.

(first entry)

04-DEC-2001 AAH99163;

BP

AAH99163/c ID AAH99163 standard; DNA; 20

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The nucleic acid sequences represent the 83P5G4 gene and the primers and adaptors used to amplify 83P5G4 DNA. 83P5G4 exhibits prostate specific expression in normal adult tissue, but it is also aberrantly expressed in many cancers including tumours of the prostate, testis, bladder, kidney, brain, bone, cervix, uterus, ovary, breast, pancreas, stomach, rectum, liver, colon and lung The 83P5G4 polynucleotide, its related protein and also peptide fragments of the protein are therefore useful for diagnosing and treating cancer. A vector comprising a polynucleotide which encodes a single chain monoclonal antibody, that immunospecifically binds to an single chain monoclonal antibody, that immunospecifically binds to an single-related protein, and a ribozyme capable of cleaving a polynucleotide having the 83P5G4 cading sequence, are both useful in the preparation of a composition for treating a patient with a cancer that cyrpesses 83P5G4. The sequences can be used in diagnostic methods to monitor the level of 83P5G4 gene products in serum, blood, unine and tissue and to thereby detect the presence of cancerous cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 55; 112pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                373 TCCTGGACCGCGACG 390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 m
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hes 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cancer
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Matches
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An isolated 83P5G4-related protein useful as a diagnostic and/or therapeutic agent in multiple cancers such as prostate, bladder and bone

Levin E;

Hubert RS, Afar DEH, Challita-Eid PM, Faris M, Mitchell SC, Jakobovits A;

WPI; 2001-514669/56.

(UROG-) UROGENESYS INC.

09-FEB-2001; 2001WO-US004426. 09-FEB-2000; 2000US-0181261P.

WO200159115-A2. Homo sapiens

16-AUG-2001.

*tag=

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modified_base
                                                                                                                                                                                                                                                                                                                       Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                        formation
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                                                                                                                                                                                                                                                                                                                                        The present invention relates to antisense compounds up to 30 nucleobases in length targeted to a E2F transcription factor 1 The invention is useful for inhibiting the expression of E2F transcription factor 1 in cells or tissues. The antisense oligomucleotides may also be used as a research agent and to prevent infection, inflammation or tumours
                                                                                                          Antisense; E2F transcription factor 1; human; infection; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human PKA C-alpha chimeric antisense oligonucleotide (ISIS# 102684).
                                                                                                                                                                                                                                                                                   Antisense compound capable of inhibiting the expression of E2F transcription factor 1, useful for preventing or delaying infection, inflammation or tumor formation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; protein kinase A; PKA catalytic subunit C-alpha inhibitor; therapy; infection; inflammation; tumour; prophylaxis; antisense; phosphorothioate backbone; chimeric; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                        Human E2F transcription factor 1 antisense oligonucleotide #71.
                                                                                                                                                                                                                                                                                                                                                                                                              3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.88+02; ative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 3 A; 9 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                 Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                        Example 15; Col 43; 40pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                    297 AAGGACCTGAGCCCCGGG 314
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            658/c
AAD09658 standard; DNA; 20 BP.
                                  AAF91365 standard; DNA; 20 BP
                                                                                                                                                                                          02-MAR-2000; 2000US-00517584.
                                                                                                                                                                                                            02-MAR-2000; 2000US-00517584
                                                                                                                                                                                                                                                Popoff I, Brown-Driver VL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAGGAACTGAGGCCTGGG
                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-SEP-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 83.3
les 15, Conservative
                                                                                                                                                                                                                            (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                 WPI; 2001-190981/19.
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Synthetic.
                                                                                                                                     Homo sapiens.
                                                                                                                                                      US6187587-B1
                                                                      04-MAY-2001
                                                                                                                                                                         13-FEB-2001.
                                                                                                                   tumour; ss
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                                                   AAF91365;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antisense oligonucleotides for inhibiting the expression of the human protein kinase A catalytic subunit C-alpha, particularly useful for preventing, delaying or treating infection, inflammation or tumor
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                      note = "Phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 1 A; 8 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mod_base= OTHER
/note= "Methoxyethyl residues"
                                                                                                         /mod_base= OTHER
/note= "Methoxyethyl residues"
                                                                                                                                                                                                                                                                                                                                                  '*tag= e
'note= "Central gap region"
/mod_base= OTHER
                                                                                                   base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                        base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Col 45; 35pp; English.
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                                                                                                                                                                                     /*tag= c
/mod_base= m5c
                                                                                                                                                                                                                                                                                                 base= m5c
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/mod_base= m5c
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                                                                                                                                                                                                                                                                  *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-407321/43.
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                                                                                                                                                                                                                                                                                                                        misc_feature
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systemic lupus erythematosus; allograft rejection; ISIS 107231; ss

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel recombinant nucleic acids useful for diagnosing, prognosing and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  treating cancer and neurological disorders, corresponds to a protein binding to alpha-catenin protein and with signal transduction function.
                                                                                                                                                                   Human; ANC_2H01 protein; catenin-binding protein; signal transduction; gene regulation; time finger protein; alphan-catenin; drug screening; therapy; cancer; neurological disorder; cytostatic; neuroprotective; PCR primer; RACE; rapid amplification of DNN end; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Integrin alpha 4; antisense; very late antigen 4; VLA4; utolmmune disease; inflammatory disease; rheumatoid arthritis; multiple sclerosis; tumour metastasis; melanoma; asthma; psoriasis; allergy; Grave's disease; Hashimoto's thyroiditis; oligonucleotide;
                                                                                                                                  Human ANC_2H01 cDNA amplifying reverse 5' RACE PCR primer, FVR359R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antisense oligonucleotide for human integrin alpha 4, ISIS 107231.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                       (VLAA-) VLAAMS INTERUNIVERSITAIR INST BIOTECHNOG.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.1%; Score 13.2; DB 1;
83.3%; Pred. No. 3.8e+02;
tive 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Van Roy F, Vanlandschoot A, Janssens B;
Example; Page 66; 160pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                286 CCAAGCTGGTGAAGGACC 303
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 350
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The sequence is an antisense oligonucleotide targetting human integrin 4, a procled in autoimmune and inflammatory diseases. The invention relates to antisense inhibitors of integrin alpha 4 which target and inhibit expression of integrin alpha 4 which target and inhibit in the expression of integrin alpha 4 muman cells or tissues, treating an animal having a disease or condition associated with expression of integrin alpha4, e.g., inflammatory disease or condition, autoimmune disease or condition including rheumatod arthritis, militiple solerosis and tumour metastases, melanoma, asthma, psoriasis, alargy, drave's disease, Hashimoto's thyroiditis, systemic lupus erythematosus and allograft rejection, and diseases or conditions enzarerised by leukocyte migration into affected tissues, preferably central nervous system tissues. The antisense molecules are also useful for reducing the reducing the adherence of cells of a first type e.g., melanoma cells or implocytes, to cells of a second type e.g., endothelial cells, by the system tissues are also endothelial cells, by the system transfer of the second type e.g., endothelial cells, by the system transfer of the second type e.g., endothelial cells, by the system transfer of the second type e.g., endothelial cells, by the system transfer of the second type e.g., endothelial cells of a first type e.g., endothelial cells of e.g., endothelial cells of e.g., endothelial cells of e.
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                                                                                                                                                                                                                                                                                                  note= "Other= All cytisines are 5-methyl cytosines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             lymphocytes, to cells of a second type e.g., endothelial cells, by inhibiting integrin alpha4 expression and thus decreasing adhesion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13...20

13...20

/*teg= 6 / mod base= OTHER / note= "Other= 2' methoxyethoxy residues"
                                                                                                                                                        *tag= a
|mod_base= OTHER
|note= "Other= Phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                          methoxyethoxy residues"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Composition for treating inflammatory and autoimmune antisense compound targeted to nucleic acid molecule alpha4 and inhibit expression of integrin alpha4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 2 A; 5 C; 12 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             /mod_base= OTHER
/note= "Other= 2' deoxy residues"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cowsert LM;
                                                                                                              cocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 32; Col 49; 49pp; English.
                                                                                                                                                                                                                                                                                                                                                                  mod_base= OTHER
                                                                                                                                                                                                                                                                            mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99US-00377309.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-450381/48.
                                                                                                              Key
modified_base
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                                            Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-OCT-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-JUL-2001
                                                                    Synthetic.
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AAS42202 standard; DNA; 20

AAS42202;

17-DEC-2001 (first entry)

Human prostate-related gene 103P2D6 cDNA nested primer #2.

103P2D6; PCR primer; DNA adaptor; prostate; testis; foetal tissue; ss; tumour; cancer; bone; ovary; breast; panoreas; colon; lung; cyrostatic; gene therapy; antibody therapy; ribozyme; serum; blood; urine; bladder; single chain monoclonal antibody; cervix; human.

Ното варіелв.

WO200162925-A2

30-AUG-2001

26-FEB-2001; 2001WO-US005996.

24-FEB-2000; 2000US-0184558P. 13-JUL-2000; 2000US-0218856P.

(UROG-) UROGENESYS INC

Raitano AB, Afar DEH, Rastegar GS, Mitchell SC, Hubert RS; Challita-Eid PM, Faris M, Jakobovits A;

WPI; 2001-557705/62.

New polynucleotide for treating and diagnosing prostate cancer is the 103P2D6 gene which encodes for 103P2D6-related proteins.

Example 1; Page 55; 132pp; English.

Sequences AAS42193-AAS42208 represent the 101P2D6 gene and the primers and adaptors used to amplify 101P2D6 DNA. 101P2D6 is not expressed in cormal adult tissue but is aberrantly expressed in some foreial tissues and many cancers including tumours of the prostate, testis, bladder, bone, cervix, ovary, breast, pancreas, colon and lung. The 103P2D6 polymuclectide, its related protein and also peptide fragments of the protein are therefore useful for diagnosing and treating cancer. A vector comprising a polymuclectide which encodes a single chain monoclonal antibody, that immunospecifically binds to an 103P2D6. The sequence, are both useful in the preparation of a composition for treating a patient with a cancer that expresses 103P2D6. The sequences can be used in diagnostic methods to monitor the level of 103P2D6 can be used in diagnostic methods to monitor the level of 103P2D6 gene products in serum, blood, unine and tissue and to thereby detect the presence of cancerous cells 

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

Gaps ö 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; ive 0; Mismatches 3; Indels Query Match Best Local Similarity 83.3 Matches 15; Conservative

373 TCCTGGACCGCGACG 390 ო 20 recressicaciós accade

8

RESULT 352

AAD19416 standard; DNA; 20 BP. AAD19416 ID AAD1 XX

(first entry) 18-DEC-2001 AAD19416; 

Human delta-6-desaturase (hD6D-1) amplifying PCR primer #1.

Delta-6-desaturase gene; D6D; lipid metabolism disorder; atopic eczema; mastalgia; rheumatoid arthritis; Sjogren's syndrome; viral infection; gastrointestinal disorder; post viral fatigue; pre-menstrual syndrome; endometriosis; cystic fibrosis; alcoholism; Alzheimer's syndrome; cardiovascular disease; trohn's disease; congenital liver disease; schizophrenia; diabetic neuropathy; nephropathy; retinopathy; cancer; arterial hypertension; atherosclerosis; chronic inflammatory disorder; autoimmune disorder; hyperholesterolaemia; atopic disorder; hb6D-1; gene therapy; human; PCR primer; ss.

Homo sapiens.

WO200170993-A2

27-SEP-2001.

26-MAR-2001; 2001WO-CA000398.

24-MAR-2000; 2000CA-02301158.

(SCOT-) SCOTIA HOLDINGS PLC.

De Antueno RJ; Allen SJ, Ponton A, Smith HL, Winther MD,

WPI; 2001-611507/70.

Nucleic acid encoding delta-6-desaturase gene useful for treating atopic eczema, mastalgia, rheumatoid arthritis, Sjogren's syndrome, gastrointestinal disorders, viral infections and post viral fatigue.

Example 4; Page 69; 164pp; English.

The invention relates to polynucleotides that control delta-6 desaturase genes (D6D) and methods useful for identifying compounds which inhibit or promote the activity of mammalian D6D. Compounds which inhibit or segments are useful for treating lipid metabolism disorders e.g. atopic eczema, mastalgia, rheumatoid arthritis, Sjogren's syndrome, gastrointestinal disorders, viral infections and post viral fatigue, premenstrual syndrome, endometriosis, cystic fibrosis, alcoholism, Alzhaimer's syndrome, candiovascular disease, Crohn's disease, cancer, companital livar disease, schizophrenia, diabetes and diabetic complications including diabetic neuropathy, nephropathy and retinopathy. Compounds of the invention are also useful for inhibiting progressive and acute disorders such as arterial hyperthension, atherosolerosis, dhronic inflammatory and autoimmune disorders, hypercholesterolaemia and other atopic disorders. D6D genes are useful in gene therapy. The present acopic else are useful in gene therapy. The present sednence

Sequence 20 BP; 6 A; 4 C; 7 G; 3 T; 0 U; 0 Other;

Gaps ö 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; rative 0; Mismatches 3; Indels 15; Conservative Query Match Best Local Similarity Matches

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RESULT 353

AAD07091/C ID AAD07091 standard; DNA; 20 BP XX AC AAD07091; XX DT 06-AUG-2001 (first entry)

The present sequence is nested primer (NP2) which is used to isolate the human six transmembrane epithelial antigen of the prostate (STEAP) CDNA fragment generated from suppression subtractive hybridisation (SSE).

STEAP is a member of cell surface serpentine transmembrane antigens.

STEAP gene is used in gene therapy. Inhibiting the development or progression of a cancer (eg. prostate, colon, bladder, lung, ovarian and pancreatic) expressing STEAP or inhibiting growth or killing cells expressing STEAP, or inhibiting growth or killing cells expresses STEAP, or inhibiting growth or killing cells expressing to the patient. Treating a patient with a cancer that composition to the patient of the patient or killing cells expressing STEAP, or inhibiting growth or killing cells expressing single chain monoclonal antibody that specifically binds to STEAP, comprises administering to the patient a vector encoding single chain monoclonal antibody coding sequence to the cancer cells and the encoded single chain monoclonal antibody is expressed intracellularly New STEAP (six transmembrane epithelial antigen of the prostate) proteins, expressed in human cancers, useful for detecting and treating NP2 primer used in isolation of STEAP cDNA fragment generated from SSH Human, cytostatic, antiproliferative, vaccine, gene therapy, six transmembrane epithelial antigen of the prostate-1, STEAP-1, prostate; colon, bladder; lung, ovarian; pancreatic, PCR primer; Raitano AB, Saffran DC, Mitchell SC; Example 1; Page 70; 187pp; English. 06-DEC-1999; 99US-00455486. 36-DEC-2000; 2000WO-US033040. Hubert RS, Jakobovits A; UROG-) UROGENESYS INC WPI; 2001-367804/38. WO200140276-A2. Homo sapiens. Afar DEH, Faris M, 

0; Gaps Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

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RESULT 354 AAS11672/c ID AAS11672 standard; DNA; 20 AAS11672;

Prostate and testis-related gene 84P2A9 cDNA nested primer #2. 24-OCT-2001 (first entry)

84P2A9; PCR primer; DNA adaptor; prostate; testis; tissue; cancer; 88; leuksemia; tumour; kidney; brain; bone; skin; ovary; breast; pancreas; colon; lung; cytostatic; gene therapy; antibody therapy; ribozyme; single chain monoclonal antibody; serum; blood; urine.

The mucleic acid sequences represent the 84P2A9 gene and the primers and adaptors used to amplify 84P2A9 DNA. 84P2A9 exhibits prostate and testis specific expression in normal adult tissue, but it is also aberrantly expressed in many cancers including leukaemia and tumnours of the prostate, testis, kidney, brain, bone, skin, ovary, breast, pancreas, colon and lung. The 84P2A9 polymuclectide, its related protein and also peptide fragments of the protein are therefore useful for diagnosing and treating cancer. A vector comprising a polymuclectide which encodes a single chain monoclonal antibody, that immunospecifically binds to an elapaba-related protein, and a ribozyme capable of cleaving a polymuclectide having the 84P2A9 coding sequence, are both useful in the preparation of a composition for treating a patient with a cancer that expresses 84P2A9. The sequences can be used in diagnostic methods to monitor the level of 84P2A9 gene products in serum, blood, urine and tissue and to thereby detect the presence of cancerous cells Levin E, Mitchell SC; New 84P2A9 gene and its encoded protein, useful for diagnosing and treating cancer, e.g. leukemia and cancer of the prostate, testis, kidney, brain or bone, or for eliciting an immune response. Challita-Eid PM, Example 1; Page 71; 149pp; English. 26-JAN-2001; 2001WO-US002651 26-JAN-2000; 2000US-0178560P Jakobovits A, Afar DEH, Hubert RS; (UROG-) UROGENESYS INC. WPI; 2001-502631/55. WO200155391-A2 02-AUG-2001 

Gaps Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels 0; 373 TCCTGGACCGCGACGACG 390 20 rechedecededadeade 3 ਨੇ

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ABL50419 standard; DNA; 20 BP 17-JUN-2002 (first entry) ABL50419; RESULT 355 ABL50419, 

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Human 158P1F4 gene nested primer (NP)2 SEQ ID NO:736.

40200216598-A2

Homo sapiens

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

Human; 158P1H4; chromosome 8q220q23, 158P1F4; chromosome 8q23; cancer; bladder cancer; immune response; cytotoxic T lymphocyte; CTL; HLA; human leukocyte antigen; helper T lymphocyte; HTL; PCR primer; adapter;

22-AUG-2001; 2001WO-US026411.

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Monitoring 158P1H4 gene products in biological sample from patient who has or is suspected of having cancer, useful for treating cancer, comprises identifying presence of aberrant 158P1H4 gene products in biological sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 45; Page 116; 209pp; English.
                                                                                                                                                                 Challita-Eid PM, Hubert RS,
Faris M, Ge W, Jakobovits A;
22-AUG-2000; 2000US-0227098P.
10-APR-2001; 2001US-0282739P.
                                                                                                                                                                                                                                                                          WPI; 2002-269357/31.
                                                                                                    (AGEN-) AGENSYS INC.
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The present invention describes a method for monitoring 158P1H4 gene products in a biological sample from a patient who has or is suspected of paving cancer. The method comprises determining the status of 158P1H4 gene products in a tissue sample from an individual, comparing the status of comparing the presence of abbrrant 158P1H4 gene products in a normal sample, and to the status of 158P1H4 gene products in a normal sample, and comparing the presence of abbrrant 158P1H4 gene products in a normal sample, 158P1H4 sequences have cytostatic activity and can be used in vaccine production. 158P1H4 polynucleotides may be used in monitoring genetic abnormalities. The 158P1H4 proteins may be used in monitoring genetic composition of the malignant phenotype, in generating and characterising concidenting the malignant phenotype, in generating and characterising concidenting the malignant phenotype, in generating and characterising concidenting the products in normal versus cancerous tissues and so elucidating the malignant phenotype, in generating and characterising cancer vaccines. Antibodies against 158P1H4 are useful in diagnostic and cytotoxic T lymphocyte (CTL) or helper T lymphocyte (HTL) responses, and simmunological reagence for detecting 158P1H4 expressing cells. The prognostic assays, and imaging methodologies. The 158P1H4 gene has been contacted to chromosome 8q22-q23, and the 158P1H4 gene also described in the present invention as represent sequences used in the present invention contents in sequences used in the present invention and produced to chromosome 8q23 and the 158P1H4 gene also described in the present invention and produces are also described in the present invention.

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

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0; Gaps
Ouery Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels
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373 TCCTGGACCGCGACGACG 390 TCCTCGGCGCGACCACG 3 20 셤 ઠે

ABL50407 standard; DNA; 20 BP ABL50407; RESULT 356 ABL50407/c 

(first entry) 17-JUN-2002 Human 158P1H4 gene nested primer (NP)2 SEQ ID NO:724.

Human; 158P1H4; chromosome 8q220q23, 158P1F4; chromosome 8q23; cancer; bladder cancer; immune response; cytotoxic T lymphocyte; CTL; HLA; human leukocyte antigen; helper T lymphocyte; HTL; PCR primer; adapter;

Homo sapiens. Synthetic. WO200216598-A2

22-AUG-2001; 2001WO-US026411 28-FEB-2002 

22-AUG-2000; 2000US-0227098P. 10-APR-2001; 2001US-0282739P.

(AGEN-) AGENSYS INC

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Levin

Raitano AB, Afar DEH,

Levin E; Raitano AB, Afar DEH, Challita-Eid PM, Hubert RS, F Faris M, Ge W, Jakobovits A;

WPI; 2002-269357/31.

Monitoring 158P1H4 gene products in biological sample from patient who has or is suspected of having cancer, useful for treating cancer, comprises identifying presence of aberrant 158P1H4 gene products in biological sample.

Example 1; Page 69; 209pp; English.

The present invention describes a method for monitoring 158P1H4 gene products in a biological sample from a patient who has or is suspected of having cancer. The method comprises determining the status of 158P1H4 gene products in a tissue sample from an individual, comparing the status of 158P1H4 gene products in a normal sample, and to the status of 158P1H4 gene products in a normal sample, and control of aberrant 158P1H4 gene products in the sample. IsBP1H4 sequences have cytostatic activity and can be used in vaccine production. 158P1H4 polynucleotides may be used in monitoring genetic abnormalities. The 158P1H4 proteins may be used in monitoring genetic abnormalities. The 158P1H4 proteins may be used in monitoring genetic abnormalities in the 158P1H4 proteins may be used in monitoring genetic abnormalities in the proteins may be used in assessing the status of 158P1H4 gene products in normal versus cancerous tissues and so elucidating the malignant phenotype, in generating and characterising concert that bind to 158P1H4 or its particular domain, and for generating cancer vaccines. Antibodies against 158P1H4 are useful in diagnostic and cytotoxic Tlymphocyte (CTL) or helper Tlymphocyte (HTL) responses, and imaging mading methodologies. The antibodies are particularly useful in bladder cancer diagnostic and prognostic assays, and imaging methodologies. The 158P1H4 gene has been correct concert diagnostic and prognostic assays, and imaging methodologies. The 158P1H4 gene has been concert to chromosome 8922-923, and the 158P1H4 gene also described in the present invention has been located to chromosome 8922-923, and the 158P1H4 gene also described in the present invention has been located to chromosome 8922-923, and the corrections when the present sequences used in the cycen prognostic assays.

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

Gaps ö 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; ive 0; Mismatches 3; Indels Query Match Best Local Similarity 83.3 Matches 15; Conservative

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AAS96899 standard; DNA; 20 BP. RESULT 357 AAS96899 

AAS96899;

(first entry) 26-FEB-2002

Human STAT3 antisense phosphorothioate oligodeoxynucleotide #106.

sTAT3; human; signal transducer and activator of transcription; ss; STAT; antisense gene therapy; Fas-mediated apoptosis; inflammatory disease; autoimmune disease; rheumatoid arthritis; cancer; breast; prostate; head; neck; brain; leukaemia; myeloma; melanoma; lymphoma; apoptosis; antiinflammatory; immunosuppressive; antirheumatic; antiarthritic; cytostatic.

```
The invention relates to antisense compounds targeted to a nucleic acid molecule encoding a signal transducer and activator of transcription (GTAT) protein, specifically STAT3, where the antisense compounds inhibit the expression of STAT3. The antisense squences are useful for inhibiting the expression of STAT3 in cells or tissues, inducing Fas-mediated approasis in cells, and sensitising cells to apoptosis. They are also useful for treating an animal having a disease or condition associated with STAT3. These disorders include inflammatory or autoimmune breast, prostate, brain and head and neck and leukaemias, myelomas, melanomas and lymphomas. Also treatable are human diseases or conditions characterised by a reduction in apoptosis or an insensitivity to apoptotic signals. The sequences of the invention can be used in clinical research, for detecting and determining the role of STAT3 in various cell functions and physiological processes and for diagnosing conditions canced with the expression of STAT3. The sequences represent cDNA encoding human STAT3 and human STAT3 oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                   Novel antisense compound useful for treating and diagnosing inflammatory diseases and cancers, is targeted to a nucleic acid molecule encoding signal transducer and activator of transcription proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 12; Page 18; 21pp; English
                                                                                                                                                                                    08-APR-1999; 99US-00288461.
06-APR-2000; 2000WO-US009054.
                                                                                                                                            11-JAN-2001; 2001US-0075881
                                                                                                                                                                                                                                                                                                                                WPI; 2002-009991/01.
                                                                                                                                                                                                                                               (KARR/) KARRAS J G.
                                                             US2001029250-A1.
  Homo sapiens
                                                                                                    11-OCT-2001
                        Synthetic.
                                                                                                                                                                                                                                                                                        Karras JG;
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0; Gaps 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; tive 0; Mismatches 3; Indels Sequence 20 BP; 2 A; 6 C; 8 G; 4 T; 0 U; 0 Other; Query Match Best Local Similarity 83.33 Matches 15, Conservative

136 CCCGCCTGCCGGTGGAGG 153 2 cccecrrecrecrecace 19 à

AAS96900 standard; DNA; 20 BP RESULT 358 g

(first entry) 26-FEB-2002 AAS96900;

Human STAT3 antisense phosphorothioate oligodeoxynucleotide #107.

STAT3; human; signal transducer and activator of transcription; ss; STAT; antisense gene therapy; Fas-mediated apoptosis; inflammatory disease; autoimmune disease; rheumatoid arthritis; cancer; breat; prostate; head; neck; brain; leukaemia; myeloma; melanoma; lymphoma; apoptosis; antiinflammatory; immunosuppressive; antirheumatic; antiarthritic; cytostatic. AAS96900

ID AAS9

X AC AAS9

XX XX ZC-E

XX XX ZC-E

XX XX ZC-E

XX XX ZC-E

XX ZC-

Homo sapiens. Synthetic.

US2001029250-A1 11-OCT-2001 

11-JAN-2001; 2001US-00758881 08-APR-1999; 99US-00288461. 06-APR-2000; 2000WO-US009054.

(KARR/) KARRAS J G.

Karras JG;

WPI; 2002-009991/01

Novel antisense compound useful for treating and diagnosing inflammatory diseases and cancers, is targeted to a nucleic acid molecule encoding signal transducer and activator of transcription proteins.

Example 12; Page 18; 21pp; English

The invention relates to antisense compounds targeted to a nucleic acid molecule encoding a signal transducer and activator of transcription (STAMI) protein, specifically STAMI, where the antisense compounds inhibit the expression of STAMI: The antisense squences are useful for the expression of STAMI in cells or tissues, inducing Fasce also useful for associated approtosis in animal having a disease or condition associated with STAMI: These disorders include inflammatory or autoimmune of disease, particularly rheumatorid arthritis, cancers, such as those of the breast, prostate, brain and head and neck and leukaemias, myslomas. Contacterised by a reduction in apoptosis or an insensitivity to cappitude appropriate signals. The sequences of the invention can be used in clinical research, for detecting and determining the role of STAMI in various cell functions and physiological processes and for diagnosing conditions associated with the expression of STAMI: The sequences represent CDNA encoding human STAMI and human STAMI oligonucleotides

Sequence 20 BP; 1 A; 7 C; 8 G; 4 T; 0 U; 0 Other;

0; Gaps 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; varive 0; Mismatches 3; Indels Query Match
Best Local Similarity 83.3
Matches 15; Conservative

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136 CCCGCCTGGCGGTGGAGG 153 3 cccecrretretretres

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AAS96833 standard; DNA; 20 BP. 26-FEB-2002 (first entry) AAS96833; AAS96833/ 

RESULT 359

Human STAT3 antisense phosphorothioate oligodeoxynucleotide #66.

STAT3; human; signal transducer and activator of transcription; ss; STAT; antisense gene therapy; Fas-mediated apoptosis; inflammatory disease; autoimmune disease; thematoid arthritis; cancer; breast; prostate; head; neck; brain; leuksemia; myeloma; melanoma; lymphoma; apoptosis; antiinflammatory; immunosuppressive; antirheumatic; antiarthritic; cytostatic

Homo sapiens. Synthetic. US2001029250-A1.

11-OCT-2001

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08-APR-1999; 99US-00288461.
  11-JAN-2001; 2001US-00758881
                        WPI; 2002-009991/01.
              (KARR/) KARRAS J G.
                   Karras JG;
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The invention relates to antisense compounds targeted to a nucleic acid molecule encoding a signal transducer and activator of transcription (STAT) protein, specifically STAT3, where the antisense compounds inhibit the expression of STAT3; me antisense sequences are useful for the expression of STAT3 in cells or tissues, inducing Fascelated apoptosis in cells, and sensitising cells to apoptosis. They are also useful for treating an animal having a disease or condition associated with STAT3. These disorders include inflammatory or autoimmune classes, particularly rheumatoid arthritis, cancers, such as those of the breast, prostate, brain and head and neck and leukaemias, myelomas. Compared to apoptocic signals. The sequences of the invention can be used in clinical capoptocic signals. The sequences of the invention can be used in clinical functions and physiological processes and for diagnosing conditions associated with the expression of STAT3. The sequences represent cDNA cascociated with the expression of STAT3. The sequences represent cDNA cascociated with the expression of STAT3 oligonuclectides
Novel antisense compound useful for treating and diagnosing inflammatory diseases and cancers, is targeted to a nucleic acid molecule encoding signal transducer and activator of transcription proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 4 A; 7 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                 Example 2; Page 13; 21pp; English
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                                Gaps
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3.1%; Score 13.2; DB 1; Length 20;
83.3%; Pred. No. 3.8e+02;
iive 0; Mismatches 3; Indels
   Query Match
Best Local Similarity 83.3°
Matches 15; Conservative
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AAS96813 standard; DNA; 20 BP. AAS96813; RESULT 360 4AS968 

(first entry) 26-FEB-2002 Human STAT3 antisense phosphorothioate oligodeoxynucleotide #46.

STAT3; human; signal transducer and activator of transcription; se; STAT; antisense gene therapy; Fas-mediated apoptosis; inflammatory disease; autoimmune disease; rheumatoid arthritis; cancer; breat; prostate; head; neck; brain; leuksemia; myeloma; melanoma; lymphoma; apoptosis; antiinflammatory; immunosuppressive; antirheumatic; antiarthritic; cytostatic.

Homo sapiens. Synthetic.

US2001029250-A1.

11-OCT-2001.

11-JAN-2001; 2001US-00758881

08-APR-1999; 99US-00288461. 06-APR-2000; 2000WO-US009054.

(KARR/) KARRAS J G.

Karras JG;

WPI; 2002-009991/01

Novel antisense compound useful for treating and diagnosing inflammatory diseases and cancers, is targeted to a nucleic acid molecule encoding signal transducer and activator of transcription proteins.

Example 2; Page 13; 21pp; English.

The invention relates to antisense compounds targeted to a nucleic acid molecule encoding a signal transducer and activator of transcription (STAT) by protein, specifically STAT3, where the antisense compounds inhibit the expression of STAT3. The antisense squences are useful for the expression of STAT3 in cells or tissues, inducing Fascalated apoptosis in cells, and sensitising cells to apoptosis. They are also useful for treating an animal having a disease or condition associated with STAT3. These disorders include inflammatory or autoimmune of disease, particularly theumatorid arthritis, cancers, such as those of the breast, prostate, brain and head and neck and leukaemias, myelomas, contacterised by a reduction in apoptosis or an insensitivity to apoptotic signals. The sequences of the invention can be used in clinical cappatic signals. The sequences of the invention can be used in clinical functions and physiological processes and for diagnosing conditions canceding human STAT3 and human STAT3. The sequences represent cDNA encoding human STAT3 and insensited with the expression of STAT3. The sequences represent cDNA encoding human STAT3 aligonucleotides 

Sequence 20 BP; 2 A; 5 C; 9 G; 4 T; 0 U; 0 Other;

0; Gaps 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; tive 0; Mismatches 3; Indel8 Query Match
Best Local Similarity 83.3
Matches 15; Conservative

1 cccccrrccrccrccacc 18

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AAS62190 standard; DNA; 20 RESULT 361 AAS62190

AAS62190;

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(first entry) 29-JAN-2002 Porcine forward PCR primer for bFGF.

Pig; muscular steatosis-modulating factor; ss; metabolic; muscular; MSMF; food supplement; obssity; hyperlipidaemia; atherosclerosis; wound healing; tumour; amyotrophic lateral sclerosis; ALS; PCR primer. HANGE STANKE STA

Sus scrofa.

WO200179287-A2.

25-OCT-2001

12-APR-2001; 2001WO-CA000509

17-APR-2000; 2000US-0197936P.

MIAC ) CANADA AGRIC & AGRI-FOOD CANADA.

Gariepy C; Pomar C, Palin M,

WPI; 2002-017600/02

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The invention relates to prognosis or diagnosis of muscular steatosis by measuring the level of a muscular steatosis modulating feator (MSMP) in a human or animal and comparing this with the level in a healthy control. Any difference indicates presence of, or predisposition to, muscular steatosis. The method is particularly used for diagnosis or prognosis of muscular steatosis. The method is particularly used for diagnosis or prognosis of conders in animal breeding. Also (ant) agonists of MSMF can be used to treat, or induce (for increasing the fat content of food) muscular study of diseases and animals. The MSMF markers are also useful in the study of diseases and animals. The MSMF markers are also useful in the atherosciences, wound healing, tumours and amyotrophic lateral sclerosis increasing the programment of the present sequence is a PCR primer used to amplify a MSMF of the invention from its gene
                         Prognosis and diagnosis of muscular steatosis, useful e.g. for selecting animals for breeding, by measuring levels of specific markers, also treating or inducing steatosis.
                                                                                                                                         Example 1; Page 39; 190pp; English.
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Sequence 20 BP; 2 A; 7 C; 6 G; 5 T; 0 U; 0 Other;

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Gaps
                           ö
Score 13.2; DB 1; Length 20;
Pred. No. 3.8e+02;
0; Mismatches 3; Indels
                                                     237 GGAGGCTGCTTCCCGGGC 254
                                                                     3 GGAGGCTTCTTCCTGCGC 20
3.1%;
                           15; Conservative
 Query Match
Best Local Similarity
Matches 15; Conserv
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ABA98342; RESULT 362 ABA98342,

BP.

55P4H4; cancer; immune response; ds; PCR primer. ABA98342 standard; DNA; 20 29-NOV-2002 (first entry) Nested primer (NP) 2. Unidentified. 

13-JUN-2001; 2001WO-US019246. 13-JUN-2000; 2000US-0211454P. (UROG-) UROGENESYS INC. 20-DEC-2001.

WO200196391-A2.

Raitano AB; Mitchell SC, Levin E, Hubert RS, Afar DEH, WPI; 2002-098053/13. Jakobovits A; Faris M,

Novel isolated 55P4H4-related protein encoded by a gene over-expressed in multiple cancers, useful as a diagnostic and/or therapeutic agent for cancer, preferably prostate cancer.

This invention relates to an isolated 55P4H4-related protein encoded by a gene that is over-expressed in multiple cancers. The polypeptide is useful for inducing an immune response to an 55P4H4 protein, providing the protein comprises of at least one T cell or B cell epitope. The immune system cell is a B cell which generates antibodies that specifically bind to the protein or is a T cell, preferably a cytotoxic T. Example 1; Page 54; 160pp; English.

cell (CTC) which kills an autologous cell that expresses the 55P4H4
protein, or a helper T cell (HTL) which secretes cytokines that
callitate the cytotoxic activity of a cytotoxic T lymphocyte. A method
is mentioned which is considered useful for monitoring the presence of
cancer in an individual, where the presence of elevated 55P4H4 mRNA or
protein expression in the test sample relative to the normal tissue
ample provides an indication of the presence or status of a cancer which
course in a prostate, kidney, testis, lung, cervix, bone, bladder, brain
ocurse in a prostate, kidney, testis, lung, cervix, bone, bladder, brain
ocurse in a prostate, kidney, testis, lung, cervix, bone, bladder, brain
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ocurse in a prostate, kidney, testis, lung, cervix, bone, bladder, brain
ocurse in a prostate, kidney, testis, lung, cervix, bone, bladder, brain
conditions associated with disregulated cell growth such as cancer and is
also useful in formal canacterized by the over-expression of 55P4H4, for
assessing the status of 55P4H4 gene products in normal versus cancerous
tissue, and to assess the presence of perturbations in specific regions
of the 55P4H4 gene. This sequence represents nested primer (NP) 2 used
during the method highlighted in the examples

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other; \$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$

ö Gaps ö 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; ive 0; Mismatches 3; Indels Query Match Best Local Similarity 83.33 Matches 15; Conservative

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ABL41837 standard; DNA; 20 BP RESULT 363 ABL41837

(first entry) 29-MAY-2002 ABL41837;

PCR primer for rat endometriotic protein ENDO-I cDNA.

Rat, endometriotic protein; ENDO-1; glycoprotein; stromal cell; endometriotic tissue; endometriosis; PCR primer; ss.

Rattus sp Synthetic

19-MAR-1998; 24-JAN-2002

US2002009718-A1.

98US-00044604. 94US-00328451. 25-OCT-1994;

WPI; 2002-215823/27. (TIMM/) TIMMS K L. Timms KL; 

Novel purified and isolated glycoprotein designated ENDO-1, useful as marker for diagnosing endometriosis in female patient suspected of having endometriosis,

Example 7; Page 10; 20pp; English.

PCR primers ABL41837-38 were used to amplify a cDNA fragment of rat endometriotic protein ENDO-1. ENDO-1 is a N-acetyl linked glycoprotein, synthesized and secreted specifically by stromal cells of endometriotic tissue origin. Human ENDO-1 has a molecular weight of 40000-55000 as determined by two-dimensional sodium dodecyl sulphate-polyacrylamide gel electrophoresis (SDS-PAGE), and has an isoelectric point of 4.0-5.5. ENDO-1 is useful as a marker for diagnosing endometriosis in a female patient suspected of having endometriosis. Endometriosis in a female patient be diagnosed by obtaining a sample from the patient, and detecting the

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The invention relates to a compound of 8-50 nucleobases in length
targeted to a nucleic acid encoding protein phosphatase 1B (PTP1B), where
the compound specifically hybridises with and inhibits the expression of
PTP1B (e.g. an antisense oligonucleotide). Also included are (1) a
COMPOUND of a native site on a mucleic acid encoding
COMPOUND of an active site on a mucleic acid encoding
COMPOUND of an active site on a mucleic acid encoding
COMPOUND of an active site on a mucleic acid encoding
COMPOUND of a supported of having a disease or condition associated
COMPOUND of a native side of a disease or condition associated
COMPOUND of a native side of a disease or condition associated
COMPOUND of a native side of a disease or condition associated
COMPOUND of a native side of a minimal comprising administering the compound; (5)
COMPOUND of a native side of a minimal compound of side and side active side of a minimal comprising administering the compound; and (6)
COMPOUND of a native side of a minimal compound of side of a minimal compound of STRIB in cells or tissues, to treat or
COMPOUND of the supression of PTPIB in cells or tissues, to treat or
COMPOUND of the supression of STRIB in cells or tissues, to treat or
COMPOUND of the supression of STRIB in cells or condition associated with
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presence of ENDO-I in the sample compared to non-endometriosis controls
                                                                                                                                                                                                                                                                                                                                                                                                                                   Antisense; protein phosphatase 1B; PTP1B; ss; probe; rat; type 2 diabbetes; obseity; ovarian cancer; chronic myeloid leukaemia; hyperproliferative disease; antidiabetic; anorectic; cytostatic; blood glucose; gene therapy.
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                                                                      Score 13.2; DB 1; Length 20;
Pred. No. 3.8e+02;
0; Mismatches 3; Indels
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                                      Sequence 20 BP; 4 A; 6 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                   Rat PTPB1 antisense oligonucleotide ISIS 111603.
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                                                                                                                                                    119 CAAGTACGGCATGCTGGC 136
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                                                                                                                                                                                                                                                                                    ABK85231 standard; DNA; 20 BP
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31-JUL-2000; 2000US-00629644.
                                                                            3.1%;
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Best Local Similarity 83.3'
Matches 15; Conservative
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FREIER S M.
MONIA B P.
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                                                                                                                                                                                                                                                                                                                            ABK85231;
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(BUTL/)
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The invention relates to a compound of 8-50 nucleobases in length targeted to a nucleic acid encoding protein phosphatase 1B (PTP1B), where the compound specifically hybridises with and inhibits the expression of PTP1B (e.g. an antisense oligonucleotide). Also included are (1) a compound of 8-50 nucleobase in length which specifically hybridises with an 8 nucleobase portion of an active site on a nucleic acid encoding PTP1B; (2) inhibiting the expression of FTP1B in cells or tissues comprising contacting the cells or tissues with the compound; treating an animal having or suspected of having a disease or condition associated with PTP1B comprising administering the compound; (4) decreasing blood
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pTPIB, such as type 2 diabetes, obesity, cancer (especially ovarian cancer, chronic myeloid leukaemia and hyperproliferative diseases in an animal having or suspected of having the disease or condition, and decreasing blood sugar levels or preventing or delaying the onset of an increase in blood glucose levels in an animal. The compound is also used in diagnosetics, herapeutics, prophylaxis, and in research reagents and kits. The present sequence is an antisense compound of the invention targetting rat PTPIB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Compound for inhibiting the expression of protein phosphatase 1B (PTP1B) and for treating diabetes, cancer, or obesity, comprises an antisense oligonucleotide targeted to nucleic acid encoding PTP1B.
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type 2 diabetes; obesity; ovarian cancer; chronic myeloid leukaemia;
hyperproliferative disease; antidiabetic; anorectic; cytostatic;
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                                                                                                                                                                                             Length 20;
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                                                                                                                                                                                                                                 3; Indels
                                                                                                                                                          Sequence 20 BP; 2 A; 6 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                             Score 13.2; DB 1;
Pred. No. 3.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rat PTPB1 antisense oligonucleotide ISIS 111615
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31-JUL-2000; 2000US-00629644.
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                                                                                                                                                                                                                                                                                                                                                                                           ABK85243 standard; DNA; 20
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Best Local Similarity 83.3:
Matches 15; Conservative
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WYATT J.
FREIER S M.
MONIA B P.
BUTLER M M.
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(WYAT/)
(FREI/)
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sugar levels in an animal comprising administering the compound; (5)
preventing or delaying the onset of a disease or condition associated
with PPPIB in an animal comprising administering the compound; and (6)
preventing or delaying the onset of an increase in blood glucose levels
in an animal comprising administering the compound. The compound is used
to inhibit the expression of PPTB in cells or tissues, to treat or
prevent or delay the onset of a disease or condition associated with
PPPIB, such as type 2 diabetes, obesity, cancer (especially ovarian
cancer, chronic myeloid leukaemia and hyperproliferative diseases in an
inmal having or suspected of having the disease or condition, and for
decreasing blood sugar levels or preventing or delaying the onset of an
increase in blood glucose levels in an animal. The compound is also used
in diagnostics, therapeutics, prophylaxis, and in research reagents and
kits. The present sequence is an antisense compound of the invention

Sequence 20 BP; 2 A; 8 C; 6 G; 4 T; 0 U; 0 Other;

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3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; tive 0; Mismatches 3; Indels
                                                                           270 CTGGAGCAGGCGGCACC 287
Query Match
Best Local Similarity 83.33
Matches 15, Conservative
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Gaps

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19 CTGGAGCAGGCCAGGACC 2 셤

ABK85035 standard; DNA; 20 BP. ABK85035; RESULT 3

ABK 8503

ABK 85

(first entry) 13-AUG-2002

Human PTP1B antisense oligonucleotide ISIS 107769.

Antisense; protein phosphatase 1B; PTP1B; ss; probe; human; type 2 diabetes; obesity; ovarian cancer; chronic myeloid leukaemia; hyperproliferative disease; antidiabetic; anorectic; cytostatic; blood glucose; gene therapy.

Homo sapiens.

JS2002055479-A1.

09-MAY-2002.

14-MAY-2001; 2001US-00854883.

18-JAN-2000; 2000US-00487368.

COWSERT L M.

WYATT J. FREIER S M. MONIA B P. BUTLER M M. (COWS/) C (WYAT/) W (FREI/) F (MONI/) M (BUTL/) E (MCKA/) N

MCKAY R.

Freier SM, Monia BP, Butler MM, Mckay R; Cowsert LM, Wyatt J,

WPI; 2002-462914/49.

Compound for inhibiting the expression of protein phosphatase 1B (PTP1B) and for treating diabetes, cancer, or obesity, comprises an antisense oligonucleotide targeted to nucleic acid encoding PTP1B.

Example 15; Page 23; 133pp; English.

The invention relates to a compound of 8-50 nucleobases in length targeted to a nucleic acid encoding protein phosphatase 1B (PTP1B), where the compound specifically hybridises with and inhibits the expression of

CC prplB (e.g. an antisense oligonucleotide). Also included are (1) a compound of 8-50 nucleobases in length which specifically hybridises with a molicebase portion of an active site on a nucleic acid encoding prplB; (2) inhibiting the expression of PrplB in acidis or tissues comprising contacting the cells or tissues with the compound; treating an animal having or suspected of having a disease or condition associated with PTPB comprising administering the compound; (4) decreasing blood sugar levels in an animal comprising administering the compound; (5) preventing or delaying the onset of a disease or condition associated with PTPB in an animal comprising administering the compound; and (6) preventing or delaying the onset of an increase in blood glucose levels or an animal comprising administering the compound; and (6) preventing or delaying the onset of an increase in blood glucose levels or inhibit the expression of PTPIB in cells or tissues, to treat or or prevent or delay the onset of a disease or condition associated with PTPIB, such as type 2 diabetes, obesity, cancer (especially ovarian connect or myelolid leukaemia and hyperproliferative diseases in an animal having or suspected of having the disease or condition, and for decreasing blood sugar levels or preventing or delaying the onset of an increase in blood glucose levels in an animal. The compound is also used in diagnostics, therapeutics, prophylaxis, and in research reagents and thus an animal human PTPIB

Sequence 20 BP; 2 A; 11 C; 5 G; 2 T; 0 U; 0 Other;

0; Gaps Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels (

303 CTGAGCCCCGGGGACCGC 320 

δ 셤 RESULT 367

멾 ABA03609 standard; DNA; 20

ABA03609;

(first entry) 08-FEB-2002 Nested primer 2 used for human 34P3D7 cDNA isolation.

Human, 34P3D7, cytostatic, vaccine, gene therapy, cancer, human leukocyte antigen, HLA, major histocompatibility complex; MHC, HLA A1, HLA A11; HLA A02, HLA A24; HLA A3; HLA B35; HLA B7; primer; 8s. 

Homo sapiens.

WO200159110-A2.

16-AUG-2001.

08-FEB-2001; 2001WO-US004094.

08-FEB-2000; 2000US-0181020P.

(UROG-) UROGENESYS INC.

Challita-Eid PM, Hubert RS, Levin Faris M, Afar DEH, Challit Mitchell SC, Jakobovits A;

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WPI; 2002-025689/03.

New gene designated 34P3D7, encoding a tissue-specific protein highly expressed in prostate cancer, for use as diagnostic and/or therapeutic target for cancers, and for eliciting an immune response.

Example 1, Page 53; 112pp; English.

The invention relates to a polynucleotide, designated 34P3D7, encoding 34P3D7-related protein, comprising a sequence of 2198 nucleotides fully

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defined in the specification. The presence of elevated 34P3D7 mRNA or protein expression indicates the presence of cancer occurring in prostate, bladder, kidney, brain, bone, cervical, uterine, ovarian, breast, pancreatic, scomach, colon, rectal leukocytes, liver, and lung tissue, and in melanocytes. An antibody against the 34P3D7-related polynucleotide complementary to 34P3D7 polynucleotide, or a ribozyme capable of cleaving the 34P3D7 polynucleotide, or a ribozyme capable of cleaving the 34P3D7 polynucleotide is useful for inhibiting the development of a cancer expressing 34P3D7 in a patient. The present sequence was used in an example demonstrating suppression subtractive hybridisation (SSH)-generated isolation of a cDNA fragment of the 34P3D7 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; calreticulin; antisense compound; hyperproliferative disorder; cancer; autoimmune disease; viral infection; cardiovascular disease; antisense therapy; cytostatic; immunosuppressive; virucide; antisense; phosphorothioate backbone; ss.
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                                                                                                                                                                                                                                                                                                                                                                                          3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; ative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human calreticulin antisense oligonucleotide, ISIS 109330.
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note= "2'methoxyethyl nucleotides"
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Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAD39537;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 368
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAD3953
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The invention relates to antisense compounds, compositions and methods for modulating the expression of calreticulin. The compositions comprise antisense compounds, particularly antisense oligonuclectides, targetted to nucleic acids encoding calreticulin. The antisense compound is useful for inhibiting the expression of calreticulin in human cells or tissues. It is also useful for treating a human having a disease or condition associated with calreticulin, e.g., hyperproliferative disorder e.g. cancer, autoimmune disease, viral infection or cardiovascular disease, by inhibiting expression of calreticulin. It is useful for disposities, therapeutics, prophylaxis and as research reagents and kits. It is also used in antisense therapy. The present sequence is an antisense compound targetted to human calreticulin. This sequence is used to study the antisense inhibition of calreticulin expression-phosphorothioate 2. MOE gapmer oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human, 125P5C8; cancer, cytostatic, breast cancer, prostate cancer;
bladder cancer; kidney cancer, colon cancer, ovarian cancer; PCR, primer;
                                                                                                                                                                                                                                                                                Novel antisense compound targeted to nucleic acid encoding calreticulin, useful for treating a human having disease or condition associated with calreticulin e.g. cancer, viral infection, autoimmune disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    . 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 4 A; 7 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human 125P5C8 gene PCR primer #3.
                                                                                                                                                                                                                                                                                                                                                          Claim 3; Page 83; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    342 GGCCGGCTGCTCTACAGC 359
/*tag= j
/mod_base= m5c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14-MAR-2001; 2001US-00809638.
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                                                                                                                30-OCT-2001; 2001WO-US049045
                                                                                                                                                 30-OCT-2000; 2000US-00702327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GGCAGGCCTCTCTACAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         002/c
AAL50002 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                    Cowsert LM;
                                                                                                                                                                                   (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                    WPI; 2002-479759/51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200272785-A2.
                                                40200236743-A2
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                                                                                                                                                                                                                  Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-SEP-2002.
                                                                               10-MAY-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 369
AALS0002/c
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Afar DEH, Raitano AB,

Hubert RS, Jakobovits A;

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The present invention relates to compositions comprising a substance that modulates the status of 125P5C8 or a molecule that is modulated by 125P5C8. The status of a cell that expresses 125P5C8 is modulated. The composition is useful for treating cancer, particularly prostate, bladder, kidney, colon, ovary or breast cancer. The 125P5C8 protein and/or a nucleotide sequence encoding the protein is useful for immunising a mammal against cancer. The present sequence is a PCR primer shown in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human/mouse C/EBP phosphorothioate antisense oligonucleotide, SEQ ID:41
                                                                                                              New composition comprising a substance that modulates the status of 125P5C8 gene or a molecule that is modulated by 125P5C8, useful for treating or preventing cancer that expresses or over expresses 125P5C8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human, C/BBP alpha; CCAAT/enhancer-binding protein alpha; CEBPA; transcription factor; tissue development; cellular function; prollferation, differentiation, adipocyte; energy metabolism; chondrogenic; ovulation; follicular development; hormonal metabolic regulation; paranlocyte development; hormonal metabolic regulation; granulocyte development; cancer; tumour formation; infection; inflammation; expression inhibition; antisense therapy; quantitative real-time PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                           Seguence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                     Example 1; Page 68; 274pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               373 TCCTGGACCGCGACGACG 390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20 TCCTCGGCCGCGACCACG 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABA02229 standard; DNA; 20 BP.
                              Faris M, Challita-Eid PM,
Morrison RK, Morrison K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-FEB-2002 (first entry)
                                                                                WPI; 2002-713510/77.
(AGEN-) AGENSYS INC.
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABA02229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 370
ABA02229
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/mod_bass= OTHER /note= "2'-methoxyethyl (2'-MOE) nucleotides. All 2' MOE cytosines are 5-methylcytosine"

*tag= b

1..20 /rag= a /mod base= OTHER /note= "Phosphorothioate linkages"

Location/Qualifiers

/mod_base= OTHER /note= "2'-methoxyethyl (2'-MOE) nucleotides. All 2' MOE cytosines are 5-methylcytosine"

*tag= c

16. .20

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cothe human CCAAT/enhancer-binding protein alpha (C/EBP alpha) gene, which inhibit it lits expression. The antisense oligonucleotides were designed to target different regions of the human C/EBP alpha Gene, which inhibit it lits expression. The antisense oligonucleotides were designed to target different regions of the human C/EBP alpha RNA, and were analysed for their effect on C/EBP alpha mRNA levels by quantitative cap-terme PCR. A similar investigation on mouse C/EBP alpha RNA, and was performed using a subset of the antisense oligonucleotides that were capable of hybridising to mouse C/EBP alpha mRNA. The C/EBP family of proteins are a family of transcription factors which regulate the expression of wide range of genes that control normal tissue development, callular function, of transcription factors which regulate the expression of wide range of genes that control normal differentiation. C/EBP alpha (also known as CEBPA) is primarily found in tissues involved in energy metabolism which have a capacity to metabolise lipids, cholesterol and other sterols. It is thought to be involved in the control and other sterols. It is thought to be involved in follicular development and ovulation, and is also involved in follicular development requisitation of metabolism, and in cycle arrest in the liver, in controlling glucose transporter GUTZ promoter activity, in the hormonal regulation of metabolism, and in cycle arrest the acvelopment and ovulation, of metabolism, and in cycle arrest of diagnosis, prevention and treatment of conditions associated with the anametric conditions or the condition, or the anametric conditions or the condition or the condition or conditions associated with the condition and the conditions associated with the condition and the conditions of the condition or conditions associated with the condition of the condition of the condition of the conditions associated with 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       103P3E8; PCR primer; DNA adaptor; prostate; bladder; kidney; colon; lbreast; rectum; stomach; tumour; cancer; cytostatic; gene therapy; sf antibody therapy; ribozyme; single chain monoclonal antibody; serum; blood; urine; tissue; human; chromosome 9q13-q21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3; Indels 0; Gaps
                                                                                                                                                                                                                                                                                                New antisense oligonuclectides for modulating the expression of CCAAT/Enhancer-binding proteins alpha, particularly useful for preventing, delaying or treating infection, inflammation or tumor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human cancer-related gene 103P3E8 cDNA nested primer #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 2 A; 9 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 15; Col 42; 44pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         335 CGACCAGGGCCGGCTGCT 352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CGGCCAGCCCAGCTGCT 19
                                                                                                                                                                                                Monia BP, Butler MM, Wyatt J;
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ID AAS95820 standard; DNA; 20 BP.
                                              13-JUN-2000; 2000US-00593589.
                                                                                                13-JUN-2000; 2000US-00593589
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                               (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                    WPI; 2002-040202/05.
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23-OCT-2001
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                                                                                                                                                                                                                                                                                                                                                                                      formation
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Gaps

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(UROG-) UROGENESYS INC

Afar DEH

Raitano AB, Mitchell SC,

Challita-Eid PM,

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Monitoring 103P3E8 gene products in sample from patient (suspected of) having cancer, useful for diagnosing, managing or treating cancers, e.g. prostate cancer, comprises determining presence of aberrant 103P3E8 gene
                                                                                              Example 1; Page 55; 128pp; English.
    12-APR-2001; 2001WO-US012181.
                12-APR-2000; 2000US-0196647P.
                           (UROG-) UROGENESYS INC.
                                                       WPI; 2002-061976/08.
                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                       Faris M, Cha
Jakobovits A;
                                                                                                                                                                                                                                                                                          AAS99443;
                                                                                    products.
                                                                                                                                                                                                                                                                    RESULT 372
                                                                                                                                                                                                                                                                          AAS99443/
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Human; chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome;
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3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human chromosome 1p36-35 PCR primer SEQ ID NO:690.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20 ICCICGGCCGCGACCACG 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABL43646 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BXBXBXEXSXBXBXBXBXBXBXB
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Sequences AAS95810-AAS95820 represent the 103P3E8 gene and the primers and adaptors used to amplify 103P3E8 DNA. 103P3E8 exhibits tissue specific expression in normal adult tissue, but it is also aberrantly expressed in many cancers including tummours of the prostate, bladder, kidney, colon, lung, breast, rectum and stomach. The 103P3E8 bladder, kidney, colon, lung, breast, rectum and stomach. The 103P3E8 bladder, comprising a polymucleotide protein and also peptide fragments of the protein are therefore useful for diagnosing and treating cancer. A vector comprising a polymucleotide which encodes a single chain monoclonal antibody, that immunospecifically binds to an 103P3E8 related protein, and a ribozyme capable of cleaving a polymucleotide having the 103P3E8 coding sequence, are both useful in the preparation of a composition for treating a patient with a cancer that expresses 103P3E8. The sequences can desired in diagnostic methods to monitor the level of 103P3E8 gene products in serum, blood, urine and tissue and to thereby detect the presence of cancerous cells
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human, 98P6C3; ss; homeodomain protein; vaccine; cytostatic. epitope; transgenic animal; immunogen; T cell; B cell; cytotoxic T cell; CTL; prostate cancer; bladder cancer; kidney cancer; lung cancer; bereast cancer; uterine cancer; cervical cancer; stomach cancer; rectal cancer; colon cancer; chromosome 4q11-q12; PCR primer; adapter; suppression subtractive hybridisation; SSH.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        373 TCCTGGACCGCGACGACG 390
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20-NOV-2001

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Gaps

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The invention relates to an isolated 98P7C3-related protein which is a homeodomain protein highly expressed in various cancers. Also include are homeodomain protein highly expressed in various cancers. Also include are polymucleotides encoding the protein or proteins 90% identical to 98P7C3, a pharmaceutical composition comprising the polymucleotides (including an expression vector comprising the 98P7C3 encoding polymucleotides) or a comparation or polymucleotides in a biological sample, monitoring the 98P7C3 protein or polymucleotides in a biological sample, monitoring the 98P7C3 protein or polymucleotides and a pharmaceutical composition comprising a modulator of 98P7C3. 98P7C3 protein, or T cell/B cell composition or polymucleotides and a pharmaceutical composition comprising a modulator of 98P7C3. 98P7C3 protein, or T cell/B cell composition in immunogens derived from it, are useful in inducing an immune response (in mammal) to a 98P7C3 protein. Upon contact with a cytocoxic T cell (CTL) the immunogens induce the CTL (with its helper T cell) to kill an exposit of cucoding the protein or epitope. The antibody is useful for delivering a cytocoxic agent to the antibody or its fragment that specifically binds crocaling the protein or epitope. The antibody is useful for delivering a cytocoxic agent to the antibody or its fragment that specifically binds conjugate. The modulator is useful for treating a patient with a cancer conjugate. The modulator is useful for treating a patient with a cancer conjugate. The modulator and colon cancer, bladder cancer, kidney cancer, breate cancer, breater sequence is objected on human chromosome 4q11-q12. The hubble of conjugation cancer is objected by the m
                                                                                                                                                                               New isolated 98P7C3-related homeodomain protein highly expressed in various cancers, useful in cancer vaccines and for generating immune response directed to 98P7C3 in mammal.
                                                  Faris M, Afar DEH, Levin E;
                                                                                                                                                                                                                                                                                                      Example 1; Page 53; 155pp; English
                                                  RS,
A;
                                                  Challita-Eid PM, Hubert
Mitchell SC, Jakobovits
                                                                                                                                     WPI; 2002-097642/13.
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The present invention describes a method of arraying genome clones. The method comprises: (a) clones of the genomic libraries contained in method comprises: (b) a clones of the genomic libraries contained in mittiwell plates numbered for discrimination are mixed in each of the multiwell plates; (b) a primer designed based on the chromosome marker sequence is added to the mixture to carry out an amplification reaction; (c) a signal corresponding to the marker is detected from the resultent amplified product to specify the discrimination Nos. of the multiwell containing the clones having said marker sequence; (d) the order of the maximum in the specified discrimination Nos. to array the multiwell containination Nos. are mixed clones are cultured and the call altertions; (f) the mixed clones are cultured and the call altertions; (f) the mixed clones are cultured and the call altertions are applified by using the above primer; (g) signals replates are specified from the detected result; and (i) the clones are reconstituted as the positions on the chromosome and arrayed. The constituted as the positions on the chromosome and arrayed. The constituted as the positions on the chromosome and arrayed. The constituted as the positions on the chromosome and arrayed. The constituted as the positions on the chromosome and arrayed. The constituted as the positions on the chromosome and arrayed. The constituted as the positions on the chromosome and arrayed. The constituted as the positions of the present invention are present proxymmers for human chromosome 21q22.1, which are present proxymmers for human chromosome 21q22.1, which are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 5 A; 8 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                     Claim 4; Page 18; 528pp; Japanese.
                       12-MAR-2001; 2001JP-00068285.
                                                                             10-MAR-2000; 2000JP-00066716.
                                                                                                                                      (RIKA ) RIKAGAKU KENKYUSHO.
(GENO-) GENOTEX YG.
                                                                                                                                                                                                                                                                            Arraying genome clones.
                                                                                                                                                                                                                      WPI; 2002-144136/19.
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Match 3.1%; Score 13.2; DB 1; Length 20; Local Similarity 83.3%; Pred. No. 3.8e+02; es 15; Conservative 0; Mismatches 3; Indels 357 AGCGACTTCCTCACTTTC 374 1 AACGACTTCCTCACGGTC 18 Query Match Matches g ò

Human PTP1B mRNA level inhibition antisense DNA #1. ABK37204 standard; DNA; 20 BP (first entry) 08-MAY-2002 ABK37204; RESULT 374 **ABK37204** 

Human, mouse, rat, protein tyrosine phosphatase 1B; PTP1B; ss, adipose, liver; kidney, metabolic disease; type 2 diabetes; obesity; cancer; hyperproliferative condition; blood serum; blood plasma; antidiabetic; blood glucose level; cytostatic; anorectic; antisense gene therapy; PTP1B mRNA level inhibition.

WO200210378-A2. Homo sapiens

07-FEB-2002.

30-JUL-2001; 2001WO-US023874.

31-JUL-2000; 2000US-00629644.

(ISIS-) ISIS PHARM INC.

Cowsert LM, Wyatt J, Freier SM, Monia BP, Butler MM, Mckay R;

WPI; 2002-180079/23.

Novel antisense compound useful for treating type 2 diabetes, cancer and obssity, is targeted to nucleic acid encoding human protein phosphatase 18, and hybridizes and inhibits PTP18 expression.

Example 15; Page 67; 142pp; English.

The invention relates to a compound targeted to a nucleic acid molecule encoding procein phosphatase 1B (PTP1B), which specifically hybridises with and inhibite the expression of PTP1B. The compounds of the invention are useful for inhibiting the expression of PTP1B in liver, kidney or adipose cells or tissues and for treating an animal, preferably human, having a disease or conditions, e.g. type 2 diabetes and obesity, and is associated with PTP1B, including metabolic hyperproliferative conditions such as cancer. The sequences are also useful for decreasing blood (serum or plasma) glucose levels in an animal e.g. a diabetic human or rodent, for preventing or delaying the onset of a disease or condition associated with PTP1B, and for preventing or delaying the onset of an increase in blood glucose levels. This sequence represents a PTP1B mRNA level inhibition antisense oligonucleotide of the invention 

Sequence 20 BP; 2 A; 11 C; 5 G; 2 T; 0 U; 0 Other;

ö Query Match
3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels

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303 CTGAGCCCCGGGGACCGC 320 

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'412/c ABK37412 standard; DNA; 20 BP. 08-MAY-2002 (first entry) ABK37412; ABK3 74 

RESULT 375

0; Gaps

Rat PTP1B mRNA level inhibition antisense DNA #129.

Human; mouse; rat; protein tyrosine phosphatase 1B; PTP1B; ss; adipose; liver; kidney; metabolic disease; type 2 diabetes; obesity; cancer; hyperproliferative condition; blood serum; blood plasma; antidiabetic; blood glucose level; cytostatic; ancrectic; antisense gene therapy; PTP1B mRNA level inhibition.

Rattus norvegicus.

WO200210378-A2.

07-FEB-2002.

30-JUL-2001; 2001WO-US023874. 31-JUL-2000; 2000US-00629644.

ISIS-) ISIS PHARM INC.

Cowsert LM, Wyatt J, Freier SM, Monia BP, Butler MM, WPI; 2002-180079/23. Novel antisense compound useful for treating type 2 diabetes, cancer and obesity, is targeted to nucleic acid encoding human protein phosphatase 1B, and hybridizes and inhibits PTPIB expression.

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73; 142pp; English
Claim 3; Page
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The invention relates to a compound targeted to a nucleic acid molecule according protein phosphatase 1B (PTP1B), which specifically hybridises with and inhibite the expression of PTP1B. The compounds of the invention are useful for inhibiting the expression of PTP1B in liver, kidney or adjonse cells or tissues and for treating an animal, preferably human, having a disease or condition associated with PTP1B, including metabolic diseases or conditions such as cancer. The sequences are also useful for decreasing blood (serum or plasma) glucose levels in an animal e.g. a diabetic human or rodent, for preventing or delaying the onset of an increase in blood glucose levels. This sequence represents a PTP1B mRNA level inhibition antisense oligonucleotide of the invention

Sequence 20 BP; 2 A; 8 C; 6 G; 4 T; 0 U; 0 Other;

Gaps ö Score 13.2; DB 1; Length 20; Pred. No. 3.8e+02; 3; Indels 0; Mismatches ch 1 Similarity 83.3%; 15; Conservative ( Query Match Best Local Similarity ઠ

270 CTGGAGCAGGGGGCACC 287 N

19 creckéckécckágakec

g

**ABK37400** RESULT

ABK37400 standard; DNA; 20

BP.

ABK37400;

(first entry) 08-MAY-2002

Rat PTP1B mRNA level inhibition antisense DNA #117.

Human; mouse; rat; protein tyrosine phosphatase 1B; PTP1B; ss; adipose; liver; kidney; metabolic disease; type 2 diabetes; obesity; cancer; hyperproliferative condition; blood serum; blood plasma; antidiabetic; blood glucose level; cytostatic; anorectic; antisense gene therapy; PTP1B mRNA level inhibition.

Rattus norvegicus.

WO200210378-A2

07-FEB-2002

30-JUL-2001; 2001WO-US023874.

31-JUL-2000; 2000US-00629644.

(ISIS-) ISIS PHARM INC

Butler MM, Freier SM, Monia BP, Cowsert LM, Wyatt J,

Mckay

WPI; 2002-180079/23.

Novel antisense compound useful for treating type 2 diabetes, cancer and obesity, is targeted to nucleic acid encoding human protein phosphatase 18, and hybridizes and inhibits PTP1B expression.

Example 16; Page 72; 142pp; English.

The invention relates to a compound targeted to a nucleic acid molecule encoding protein phosphatase 1B (PTP1B), which specifically hybridises with and inhibits the expression of PTP1B. The compounds of the invention are useful for inhibiting the expression of PTP1B in liver, kidray adipose calls or tissues and for treating an animal, preferably human, having a disease or condition associated with PTP1B, including metabolic

diseases or conditions, e.g. type 2 diabetes and obesity, or hyperproliferative conditions such as cancer. The sequences are also useful for decreasing blood (serum or plasma) glucose levels in an animal e.g. a diabetic human or rodent, for preventing or delaying the onset of a disease or condition associated with PTPLB, and for preventing or delaying the onset of delaying the onset of an increase in blood glucose levels. This sequence represents a PTPLB mRNA level inhibition antisense oligonucleotide of the invention 8888888888888

Sequence 20 BP; 2 A; 6 C; 8 G; 4 T; 0 U; 0 Other;

Gaps .. 0 Length 20; 3; Indels Score 13.2; DB 1; Pred. No. 3.8e+02; 0; Mismatches 3.1%; Local Similarity 83.3 Query Match Best Loca Matches

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RESULT 377

ABT12959 standard; DNA; 20 BP. ABT1295

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ABT12959;

(first entry) 17-JAN-2003 Mycobacterium tuberculosis-specific DNA sequence #46.

Mycobacterium detection method; PCR; primer; probe; ss.

Mycobacterium tuberculosis.

WO200274991-A2

26-SEP-2002.

20-MAR-2002; 2002WO-GB001308

20-MAR-2001; 2001GB-00006949

(NORC-) NORCHIP AS. (ALLA/) ALLARD S J.

Karlsen F;

WPI; 2002-750564/81.

Detecting the presence of Mycobacterium tuberculosis in a test sample, comprises inducing mRNA expression of Mycobacterium tuberculosis and detecting the induced mRNA.

Claim 8; Page 14; 70pp; English.

The invention comprises a method for detecting the presence of a microorganism (particularly Mycobacterium tuberculosis) in a test sample. The method of the invention comprises exposing the test sample to an inducer that is capable of inducing the expression of at least one gene in the micro-organism and then testing for the presence of mRNA from this gene. The method of the invention is useful for detecting an mRNA that is expressed in a species of Mycobacterium (e.g. Mycobacterium tuberculosis). The present DNA sequence represents a Mycobacterium specific nucleotide which can be used as a primer or probe in the method the invention 

Sequence 20 BP; 5 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

ö 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; tive 0; Mismatches 3; Indels 3.1 Best Local Similarity 83.3 Matches 15; Conservative

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Gaps

374 CCTGGACCGCGACGACGG 391

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2 CCTGGACTGGGACTACGG 19

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ABQ62315 standard; DNA; 20 BP.

RESULT 378

ABQ6231

16-AUG-2002 (first entry)

ABQ62315;

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Antisense PCR primer used to amplify GAPDH cDNA
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PCR; primer; ss; adenoviral vector; toxic gene; cancer; promoter; liver; gene therapy; suicide gene; gastrointestinal cancer; pancreatic cancer; hepatctropism; adenovirus; intrahepatic tumour; liver dysfunction; cyclooxygenase-2; Cox-2; carcinogenesis; colon cancer; tumour; liver toxicity; GAPDH; glyceraldehyde-3-phosphate dehydrogenase.

Example 2; Page 4; 35pp; English

The invention discloses an adenoviral vector for the selective expression of a toxin gene in a cancer cell. The toxin gene is operably linked to a promoter of a gene with undetectable expression in liver, so that expression of the toxin gene is reduced in liver cells. Adenoviral expression of the toxin gene is reduced in liver cells. Adenoviral vectors and adenoviral gene therapy have been used to introduce suicide/toxic genes to advanced gastrointestinal or pencreatic cancer cells. This technique has a problem due to the hepatotropism of the adenovirals for systemically administered adenoviral vectors localise comprised to the liver, where the suicide gene therapy of intrahepatic tumour leads to severe liver dysfunction. Cyclooxygenase-2 (Cox-2) has virtually undetectable expression in most tissues, but is closely linked to recrinogenesis and progression of colon cancers. The promoter of Cox-2 therefore has a tumour "on" liver "off" expression profile, which can be utilised in the adenoviral gene therapy vectors. The adenovirus is used to the liver dysfunction cancer cells. The adenovirus is used to satrointestinal or pancreatic cancer cells. The expression corrected the antisense PCR primer which was used to amplify glyceraldehyde-3. profile of cyclooxygenase-2 (Cox-2) cDNA in various tissues

Sequence 20 BP; 4 A; 7 C; 6 G; 3 T; 0 U; 0 Other;

The invention comprises antisense oligonucleotides designed to inhibit expression of Syntaxin 4 interacting protein. The antisense oligonucleotides of the invention are useful for inhibiting the expression of Syntaxin 4 interacting protein in cells or tissues. The antisense oligonucleotides are also useful for treating an animal having a disease or condition associated with Syntaxin 4 interacting protein (e.g. diabetes, obesity or a skeletal muscle disorder). The antisense oligonucleotides can also be used to prevent or delay infection, inflammation and tumour formation. The present DNA sequence represents a human Syntaxin 4 interacting protein antisense oligonucleotide. NoTE: The methoxyethyl wings

Novel antisense compound that hybridizes and inhibits nucleic acid molecule encoding Syntaxin 4 interacting protein, useful for treating diabetes, obesity and skeletal muscle disorder.

Monia BP, Freier SM, Wyatt JR;

WPI; 2002-404952/43.

(ISIS-) ISIS PHARM INC

19-SEP-2001; 2001WO-US029251. 22-SEP-2000; 2000US-00668313.

WO200224864-A2. Homo sapiens.

28-MAR-2002

Claim 3; Page 84; 154pp; English.

0; Gaps Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels 0

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85 CAGTGGACATCACCACGT 102 요 ઠે

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Gaps

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Query Match
3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels

116 CAGCAAGTACGGCATGCT 133

20

CAACAAGTAGTGCATGCT

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ABX14127 standard; DNA; 20

RESULT 379

ABX14127

(first entry)

27-FEB-2003

ABX14127;

SAXAXEX

Sequence 20 BP; 6 A; 4 C; 4 G; 6 T; 0 U; 0 Other;

AAL38219 standard; DNA; 20 AAL38219; 

BP.

(revised)
(first entry) 29-AUG-2003 15-AUG-2002 Human BH3 interacting domain death mRNA agonist inhibitor SEQ ID 62

An adenoviral vector containing a toxin gene under control of a promoter with undetectable expression in the liver is useful to treat gastrointestinal or pancreatic cancer with reduced liver toxicity. 07-DEC-2001; 2001US-00005964 05-DEC-2000; 2000US-0251375P Curiel DT, Yamamoto M; WPI; 2002-697880/75 (CURI/) CURIEL D T. (YAMA/) YAMAMOTO M. JS2002107219-A1. Homo sapiens 08-AUG-2002. 

Human, antisense gene therapy, Syntaxin 4 interacting protein, ss; antisense oligonucleotide; diabetes; obesity; skeletal muscle disorder; inflammation; tumour formation; phosphorothioate backbone; 2'-0-methoxyethyl wing.

Human syntaxin 4 interacting protein antisense oligonucleotide 54,

rng.res

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Hepatotrophic; immunomodulatory; cytostatic; antiinflammatory; hepatitis; haemostatic; BH3 interacting domain death agonist; liver disease; haematopoietic disorder; developmental disorder; immunological disorder; hyperproliferative disorder; apoptosis; human; chimeric; 2'-methoxyethyl; 2'-MOB; phosphorothioate backbone; ds.
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Homo sapiens

Chimeric.

WO200220547-A1.

14-MAR-2002

31-AUG-2001; 2001WO-US027316.

07-SEP-2000; 2000US-00657346. 07-MAR-2001; 2001US-00800631.

(ISIS-) ISIS PHARM INC.

Wyatt JR; Zhang H, WPI; 2002-393838/42.

Novel antisense compound targeted to nucleic acid molecule encoding the BH3 interacting domain death agonist, useful for treating animals with diseases associated with BH3 interacting domain death agonist, e.g. hepatitis.

Claim 3; Page 87; 171pp; English.

The invention relates to a compound 8 to 50 nucleotides in length
targeted to a nucleic acid molecule encoding a BH3 interacting domain
death agonist, where the compound specifically hybridises with and
inhibits the expression of the BH3 interacting domain death agonist. The
compound of the invention is useful for inhibiting the expression of the
BH3 interacting domain death agonist in cells or tissues. The compound is
also useful for treating an animal having a disease or condition
associated with the BH3 interacting domain death agonist, e.g.
cassociated with the BH3 interacting domain death agonist, e.g.
disorder, immunological disorder, or a disease or condition of the liver
cissue full for dispositios, therapeutics, prophylaxis and as research
creagents and kits. This polynucleotide sequence represents an antisense
coligonucleotide inhibitor of the DNA from human BH3 interacting domain
death agonist RNA of the invention. NOTE: This sequence is a diffmentic
coligonucleotide wings'. The wings are composed of 2' methoxyethyl (2'
"NOE) nucleotides. The internucleoside (backbone) linkages are
chosp nucleotides. The internucleoside (backbone) linkages are
chosp nucleotides. The internucleoside (backbone) linkages are
cologia to standardise OS field) 

Sequence 20 BP; 4 A; 4 C; 10 G; 2 T; 0 U; 0 Other;

Gapa ö Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels

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114 CGCAGCAAGTACGGCATG 131

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Human C/EBP beta phosphorothioate antisense oligonucleotide, SEQ ID:40.

M M M MOE Human, C/EBP beta, CCAAT/enhancer-binding protein beta, C/EPB2; LAP;
TCF5; CRP2; NFL16; IL6DBP, NF-M, AGP/EBP; Apc/EBP; transcription factor;
tissue development; cellular function; proliferation; differentiation;
hormone responsiveness; oxidative stress response;
IL-6 signalling mediator; interleukin-6; carbohydrate metabolism;
immunity; Thi response; female fertility; gluconeogenesis; ovarian;
cancer; tumour formation; type II; diabetes; infection; inflammation;
expression inhibition; phosphorothioate; antisense oligonucleotide; ss. 1..5 /*tag= b /mod = 0.7HER /note= "2'-methoxyethyl (2'-MOE) nucleotides, All 2' cytosines are 5-methylcytosine" 'n /*tag= c /*tag= CTHER //mcd_base= OTHER //note= "2'-methoxyethyl (2'-WOE) nucleotides. All cytosines are 5-methylcytosine" 1..20 *trag= a /mcd base= OTHER /note= "Phosphorothioate linkages" Location/Qualifiers 16. .20 Key modified_base modified_base modified_base Homo sapiens. 

US6271030-B1

07-AUG-2001.

14-JUN-2000; 2000US-00593711.

14-JUN-2000; 2000US-00593711.

(ISIS-) ISIS PHARM INC

Butler MM, Wyatt J; Monia BP,

WPI; 2002-214451/27.

Novel antisense compound targeted to nucleic acids encoding human or mouse CCAAT/enhancer binding protein (C/EBP) beta, useful in vitro for inhibiting expression of human or mouse C/EBP beta in cells/tissues.

Claim 1; Col 42; 69pp; English.

Sequences AB194252-AB194476 represent antisense oligonucleotides targeted to the human or mouse CCAAT/enhancer-binding protein alpha (C/BBP alpha) and which inhibit its expression. The antisense oligonucleotides were designed to target different regions of the human and/or mouse C/BBP alpha NA, and were analysed for their effect on C/BBP alpha mRNA levels alpha RNA, and were analysed for their effect on C/BBP alpha mRNA levels control tormal tissue development, cellular function, cellular control normal tissue development, cellular function, cellular control and functional differentiation and series responses control and processed and series has beta is control and its a mediator of IL-6 (incerleuke) signalling. C/BBP beta is control cound in the lung. It is also expressed at a higher conditions are useful for diagnosis, prevention and treatment of impairment of insulin secretion in type II diabetes. The oligonucleotides conditions associated with C/BBP beta expression, guch as cancer conditions associated with C/BBP beta expression, guch as cancer conditions associated with C/BBP beta expression, guch as cancer conditions or inflammation. Guabetes (particularly very Inflammation.)

4 T; 0 U; 0 Other; A; 9 C; 5 G; BP; 2 Sequence 20

Wed Apr 21 12:58:21 2004

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                           Gaps
                           ;
Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels
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Human 83P2H3 cDNA isolation nested PCR primer 2. BP 422/c ABK67422 standard; DNA; 20 02-JUL-2002 (first entry) ABK67422; RESULT 38 ABK67422/ 

Human; human leukocyte antigen; HLA; immunogen; 83P2H3; CaTrF2E11; calcium transport protein; cancer; prostate cancer; cytostatic; chromosome 7q34; chromosome 12q24.1; T cell; B cell; Bs; primer.

Homo sapiens.

WO200214361-A2.

21-FEB-2002.

17-AUG-2001; 2001WO-US025782

17-AUG-2000; 2000US-0226329P

(AGEN-) AGENSYS INC.

Afar DEH; Raitano AB, Challita-Eid PM, Faris M, Saffran DC, Levin E, Hubert RS, Ge W, Jakobovits A;

WPI; 2002-269179/31.

Monitoring 83P2H3 gene products for monitoring the presence of cancer in a subject, comprises determining the status of 83P2H3 gene products in a tissue sample from the subject and comparing it to a normal sample.

Example 1; Page 76; 270pp; English.

The invention relates to monitoring 83P2H3 (a calcium transport protein whose gene is located on chromosome 7q34) gene products in a biological eargiele from a parient who has or is suspected of having cancer cancer), comparige (a) determining the status of 83P2H3 gene products expressed by calls in a tissue sample from an individual and (b) compariging the status to the status of 83P2H3 gene products in a normal sample. Also included are modulators of 83P2H3 function or status, generating antibodies/immune response against 83P2H3 conjugating the agent to an entigen) binding peptides derived from the protein delivering a cytotoxic agent to a cell cypridom that produces the recombinant protein, a non-human transgenic animal that produces the recombinant protein, a non-human transgenic animal that produces the recombinant protein, a non-human of the anti-83P2H3 antibody, a vector comprising a non-human of the anti-83P2H3 antibody, a vector comprising a non-conformal antibody that comprises the wortable domains of the havy and inducing the presence of cancer in an individual. The modulator is monitoring the presence of cancer in an individual. The modulator is cancer that expresses 83P2H3. The immunological methods are useful for cancer that expresses 83P2H3. The immunological methods are useful for

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generating an immune response against 83P2H3, and for detecting the presence of 83P2H3-related protein or polynuclectide in a biological sample from a patient who has or who is suspected of having cancer. The antibody is useful in prostate cancer diagnosis, prognosis, imaging methodologies and treatment, to detect and quantify 83P2H3 and mutant 83P2H3-related proteins, for purifying a 83P2H3-related protein, for isolating 83P2H3 homologues/related molecules, and for generating antidiotypic antibodies that mimic the 83P2H3 protein. The present sequence is a PCR primer used in the isolation of cDNA encoding 83P2H3 or its related protein CaTrF2EH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human, cytostatic, 85P1B3; cancer; immunogen; ss; primer; PCR; chromosome 15q14.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human cDNA 85P1B3 nested PCR primer 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           373 TCCTGGACCGCGACGACG 390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABX70514/C

ID ABX70514/C

XA ABK70514;
XX ABK70514;
XX ABK70514;
XX Human; Cytostatic; B5PlB3; Car
XX Homo sapiens.
XX W0200218578-A2.
XX AGEN-102.
XX Y AGEN-102.
XX Y AGEN-102.
XX X AGEN-1 AGENSYS INC.
XX AGENSYS INC.
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XX AGEN-1 AGENSYS INC.
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XX AGEN-1 AGENSYS INC.
XX AGENSYS INC.
XX AGENSYS INC.
XX AGEN-1 AGENSYS INC.
XX AGENS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TCCTCGGCCGCGACCACG 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-AUG-2001; 2001WO-US026838.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 383
              888888888888
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The invention relates to a composition comprising a substance that modulate the status of 85P1B3, where the status of a cell expresses 85P1B3 gene product is modulated. Also included are a composition comprising a perduct is modulated. Also includes an as position, an any whole number increment up to 229 that includes an as position any whole number increment up to 229 that includes an as position. It has been a position having a value greater than 0.5 in the percent accessible residue profile, an as position having a value greater than 0.5 in the average flexibility profile; an as position having a value greater than 0.5 in the average flexibility profile; an as position having a value greater than 0.5 in the average flexibility profile; or an as position having a value greater than 0.5 in the becareful average flexibility profile; a polymucleotide that encodes analogue peptide of 8, 9, 10 or 11 contiguous residues of the 85P1B3 protein; a recombinant protein comprising the antigen-binding region of a monoclonal antibody; a non-human transgenic animal that produces an antibody that binds to the 85P1B3 protein; a Composition for modulating the status of 8571B3 protein or a molecule comprising a substance e.g. antibody specific to, nucleic acid encoding, or ribozyme of 8571B3. Example 1; Page 76; 201pp; English.

Ge W, Challita-Eid

Faris M, Hubert RS, Afar D,

hybridoma that produces antibody specific to the protein; a single chain monoclonal antibody (MAb) that comprises the variable domains of the heart and monoclonal antibodies specific to the protein; a vector comprising a polymucleotide that encodes the MAb; inhibiting growth of cancer cells or treating a patient who bears cancer cells that expresses the protein, antibody, polymucleotide encoding the protein, antibody polymucleotide cancer cells that cleaves the polymucleotide and T cells that specifically risocomize the protein; and generating a mammalian immune response directed to the protein and generating a mammalian immune system to an immunogenic portion of the protein or polymucleotide. The composition, which comprises an antibody specific to the protein, is useful for delivering a cytocoxic agent to a cell that expresses the protein by providing a cytocoxic agent to acell that expresses the protein by conjuding a cytocoxic agent to acell that expresses the protein for the mattibody and exposing the cell to the antibody-agent conjugate. The methods are useful for inhibiting crowth of cancer cells or treating a mammalian immune response expresses the protein, for detecting the presence of the protein or polymucleotide in a biological sample in a patient who has or who is suspected of having cancer and for monitoring SPIRB3 in a biological sample from a patient who has or who is sample from a patient who has or who is sample for mere do numan chromsome 15q14. The present

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

Gaps ö Query Match
3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels 373 TCCTGGACCGCGACGACG 390 ઠે

20 rccreeccecedaccace 3

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ABI93630 standard; DNA; 20 BP.

ABI93630;

(first entry) 15-FEB-2002

Capture oligonucleptide Zip ID#717 oligo #9.

Human, K-ras, PCR primer, probe, capture probe, mutation detection, ligase detection reaction; LDR, p53; BRCA1; BRCA2; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndrome, obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forensit; environmental monitoring; food industry; feed industry; se. AB193636
AB19366
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Synthetic.

WO200179548-A2.

25-OCT-2001.

04-APR-2001; 2001WO-US010958.

14-APR-2000; 2000US-0197271P.

CORR ) CORNELL RES FOUND INC.

Favis R, Gerry NP, Zirvi M, Barany F,

Kliman R;

MPI; 2002-034366/04.

Designing capture oligonucleotide probes for use on a support to which complementary oligonucleotides hybridize with little mismatch.

Example 5; Fig 29; 300pp; English.

The present invention describes a method (M1) for designing capture

cc oligonucleotide probes (I) for use on a support to which complementary cligonucleotide probes (II) will hybridise with little mismatch, where cligonucleotide probes (II) will hybridise with little mismatch, where cc (I) have melting temperatures within a narrow trange. The method is useful cc e.g. Salmonella, Listeria monocytogenes and Haemophilus influenza, fungal infectious agents cc c.g. Salmonella, Listeria monocytogenes and Haemophilus influenza, fungal infectious agents cc fifteetious agents cc rypteococcus nectormans, Candinesia albicans and cc speciallus funigautus, viruses e.g. T-cell lymphocytotrophis citus, Epstein-Barr virus and pollo virus, and parasitic infectious agents cancer from Onchoverva volvulus, Entamoeba histolytica and Dracumculus can as 1 hydroxylase deficiency, Turner Syndrome and obesity defects.

Cc alected from Onchoverva volvulus, Entamoeba histolytica and Dracumculus cancer involving oncogenes, tumour suppressor genes, or genes involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCA1 gene, buman papillomavirus types 16 and 18 and liver cancers. The method is also used for environmental monitoring, forensics and the food and feed industry, detecting comprises scanning (using e.g. a scanning electron microscope and infrared microscope) the support at the cc presence or absence of the target mucleotide sequences. ABI82074 to present oligonucleotide sequences used in the exemplification of the present oligonucleotide sequences used in the exemplification. the present invention \$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$

Sequence 20 BP; 4 A; 8 C; 5 G; 3 T; 0 U; 0 Other;

; 0 Length 20; 3; Indels Score 13.2; DB 1; Pred, No. 3.8e+02; 0; Mismatches 3; Query Match 3.1%; Best Local Similarity 83.3%; Matches 15; Conservative (

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29 GGGCTGGGACGAAGATGG 46

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RESULT 385 AB1929

ABI92926 standard; DNA; 20 BP.

AB192926;

15-FEB-2002 (first entry)

Capture oligonucleptide Zip ID#13 oligo #9.

Human, K-ras, PCR primer, probe, capture probe, mutation detection, ligase detection reaction, LDR, p53; BRCA1, BRCA2, infectious disease, infection, 21 hydroxylase deficiency, Turner Syndrome; obesity, cancer, oncogene, tumour suppressor; human papillomavirus, forenaic; environmental monitoring; food industry, feed industry; ss.

Synthetic.

WO200179548-A2 CXSXEEXSXEXEXEXEXEXEXEXEXEXEXEXCXC

25-OCT-2001.

04-APR-2001; 2001WO-US010958.

14-APR-2000; 2000US-0197271P.

FOUND INC. (CORR ) CORNELL RES

Gerry NP, Favis R, Kliman R; Barany F, Zirvi M,

WPI; 2002-034366/04.

Designing capture oligonucleotide probes for use on a support complementary oligonucleotides hybridize with little mismatch.

Example 5; Fig 29; 300pp; English.

The present invention describes a method (M1) for designing capture

cc oligonucleotide probes (I) for use on a support to which complementary cligonucleotide probes (II) will hybridise with little mismatch, where coligonucleotide probes (II) will hybridise with little mismatch, where (I) have melting temperatures within a narrow range. The method is useful cor detecting infectious diseases caused by bacterial infectious agents c.g. Salmonella, Listeria monocytogenes and Haemophilus influenza, fungal infectious agents e.g. Cryptococcus neoformans, Candida albicans and Aspergillus fumigautus, viruses e.g. T-cell lymphocytotrophis cirus, Epstein-Barr virus and polito virus, and parasitic infectious agents selected from Onchoverva volvulus, Entamocba histolyrica and Dracunculus cas 21 hydroxylase deficiency, Turner Syndrome and obseity defects.

CC as 21 hydroxylase deficiency, Turner Syndrome and obseity defects.

Detecting cancer involving oncogenes, tumour suppressor genes, or genes involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCA1 gene, cancer is specifically associated with a gene selected from BRCA1 gene, psi on used for environmental monitoring, forensics and the food and feed industry, detecting comprises scanning (using e.g. a scanning electron microscope and infrared microscope) the support at the correlating (using a computer) identified ligation to a presence or absence of the target nucleotide sequences. ABIS2074 to of the present invention

Sequence 20 BP; 3 A; 4 C; 8 G; 5 T; 0 U; 0 Other;

Gaps ö Query Match
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels

a

AAL40496 standard; DNA; 20 BP.

AAL40496;

19-SEP-2002 (first entry)

158P1D7 cDNA related PCR primer SEQ ID No 668.

Cytostatic; 158P1D7; cancer; bladder cancer; mouse; rat; rabbit; dog; cat; cow; horse; human; vaccine; gene therapy; PCR; primer; ss.

Homo sapiens.

WO200216593-A2.

28-FEB-2002.

22-AUG-2001; 2001WO-US026276.

22-AUG-2000; 2000US-0227098P. 10-APR-2001; 2001US-0282739P.

(AGEN-) AGENSYS INC.

Faris M, Hubert RS, Raitano AB, Afar DBH, Challita-Eid PM, Jakobovits A;

Levin E;

WPI; 2002-425659/45.

New compositions comprising a gene (designated 158P1D7), its encoded protein or their modulators, useful for treating or diagnosing cancers, particularly bladder cancer, in mammals (e.g. dogs, cats, cows, horses or numans) 

Example 1; Page 68; 181pp; English

The invention relates to a novel nucleic acid, designated 158P1D7. The compositions are useful for treating or diagnosing cancers, particularly bladder cancer, in mammals (e.g. mice, rats, rabbits, dogs, cats, cows, horses or humans). The compositions are also useful for monitoring genetic abnormalities and in preparing cancer vaccines. The nucleic acid of the invention can be used in gene therapy to treat the said disorders. This polynucleotide sequence represents a PCR primer of the 158F1D7 CDNA of the invention 888888888888

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels

g

RESULT :

AAL53476 standard; DNA; 20 BP. 

16-JAN-2003 (first entry)

Zinc transporter protein 108P5H8 nested primer 2.

Cytostatic; gene therapy; vaccine; zinc transporter protein 108P5H8; cancer; breast; colon; ovarian; lung; humoral; cellular immune response; passive immunisation; PCR; primer; ss.

Unidentified

WO200260953-A2.

08-AUG-2002.

17-DEC-2001; 2001WO-US049133.

15-DEC-2000; 2000US-0256210P.

(AGEN-) AGENSYS INC.

Mitchell SC; Challita-Bid PM, Faris M, Afar DEH, Hubert RS, ^N Levin E, Morrison KJM, Raitano AB, Jakobovits A,

WPI; 2002-627469/67.

Composition comprising a substance that modulates the status of a zinc transporter protein (108P5H8), useful in diagnosing and treating patients with cancer that express 108P5H8, such as breast, colon, ovarian or lung cancer.

Example 1; Page 95; 309pp; English.

The invention relates to a new composition comprising a substance that modulates the status of a zinc transporter protein, designated as modulates. The composition is 108P5H8, or a molecula that is modulated by 108P5H8. The composition is useful in diagnosing, preventing, prognosticating or treating patients with cancer that expresses 108P5H8, such as breast, colon, ovarian or lung cancer. The 108P5H8 ene or its fragment can be used to elicit a humoral or cellular immune response. The antibodies are useful in active or passive immunisation. The 108P5H8 polymucleotides are useful in active and primers for the amplification or detection of 108P5H8 genes, as coding sequences for directing the expression of 108P5H8 genes. The polymucleotides of the invention can be used to treat disorders by gene therapy. This polymucleotide sequence represents a zinc transporter protein 108P5H8 related PCR primer of the invention

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The invention relates to a novel composition comprising a substance that modulates the status of a protein, 121P2A3. The composition of the modulates the status of a protein, 121P2A3. The composition of the invention has cytostatic and immunostimulant activity, and is useful as a vaccine. The 121P2A3 proteins and polymucleotides are useful for characterising cytogenetic abnormalities of this chromosomal cours, as tools that can be used to delineate cytogenetic abnormalities of in the chromosomal region that encodes 121P2A3 that may contribute to in the chromosomal region that encodes 121P2A3 that may contribute to in normal versus cancerous tissues. The proteins are useful for generating and characterising domain-specific antibodies, for identifying agents or cellular factors that bind to 121P2A3 or a particular structure domain, and in various therapeutic and diagnostic contexts, including cancer vaccines. The antibodies or T cells reactive with the product are useful in passive or active immunisation, and in imaging methodologies for the management of cancer. The present sequence represents an nested for the management of cancer. The present sequence represents an nested CPC suppression subtractive hybridisation (SSH) reactions
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Afar DEH, Saffran D, Morrison K, Morrison RK, Ge W, Jakobovits
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; 121P2A3; cytostatic; immunostimulant; vaccine; PCR; primer; humoral immune response; ss; callular immune response; ss; suppression subtractive hybridisation; SSH.
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                                                                      Score 13.2; DB 1; Length 20;
Pred. No. 3.8e+02;
0; Mismatches 3; Indels
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Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human 121P2A3 post-SSH nested PCR primer 2.
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83.3%; Pred
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25-APR-2001; 2001US-0286630P.
22-JUN-2001; 2001US-0300373P.
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                                                              Query Match
Best Local Similarity 83.3
Matches 15; Conservative
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DB 1; Length 20;

3.1%; Score 13.2;

Query Match

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention describes a new composition comprising a substance that modulates the status of 158P3D2 or a molecule that is modulated by 158P3D2, where the status of a cell that expresses 158P3D2 is modulated. The composition is useful for treating cancer. This sequence represents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New composition comprising a substance that modulates the status of 15893D2 or a molecule that is modulated by 15893D2, useful for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human, antisense, lung dysfunction, nasal airway dysfunction;
antiinflammatory steroid; ubiquinone, antiinflammatory; antiallergic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
            Gaps
                                                                                                                                                                                                                                      158P3D2; cytostatic; gene therapy; vaccine; cancer; PCR; primer; ss.
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Pred. No. 3.88+02;
0; Mismatches 3; Indels
              Indels
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Afar DEH, Ge W, Raitano AB, Challita-Eid PM;
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  Pred. No. 3.8e+02;
); Mismatches 3;
                                                                                                                                                                                                              Novel protein 158P3D2 associated primer #4.
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                                       373 TCCTGGACCGCGACGACG 390
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25-APR-2001; 2001US-0286630P.
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 83.3%;
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                                                                                                                                 ACA64671 standard; DNA; 20
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Best Local Similarity 83.3
Matches 15; Conservative
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                15; Conservative
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Best Local Similarity
Matches 15; Conserv
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                                                                                                                                                                                                                                                                     Synthetic.
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                                                                                                          RESULT 389
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cytostatic; gene therapy;
antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
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Homo sapiens.

WO200285308-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Katz E, Pabalan J, Aguilar D; Li Y, Sandrasagra A, K Tang L, Shahabuddin S; Nyce JW, I Miller S,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 13919; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the intifiation codon, coding region, 5. or 3. end genomic flanking regions, 5. and 3. intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal alraway dysfunction and a second active agent comprising an antiinflammatory steroid and ubjquinone. A composition of the invention has antiinflammatory, antiallargic, antisthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition may have a use in antisense gene therapy. The composition may have a use in antisense gene therapy. The composition may have a cust in antiallammatory steroid in a subject, for reducing or depleting levels of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or receptor, producing bronchodilation, increasing levels of ubjquinone or lung surfactant in a subject, to receptor, producing bronchodilation, increasing levels of ubjquinone or lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pot_sequences

Sequence 20 BP; 3 A; 7 C; 9 G; 1 T; 0 U; 0 Other;

Gaps ö Query March 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels

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ABZ86355 standard; DNA; 20 BP ABZ86355; 

(first entry) 17-OCT-2003

Human oligonucleotide sequence.

Human, antisense, lung dysfunction, nasal airway dysfunction; antiinflammatory, antiallergic;

antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P

(EPIG-) EPIGENESIS PHARM INC

Aguilar D; Katz E, Pabalan J, Li Y, Sandrasagra A, Ka , Tang L, Shahabuddin S; Miller S, Nyce JW,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

claim 15, SEQ ID NO 1597; 872pp; English.

first active agent comprising an oligonuclectide antisense to the invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonuclectide antisense to the initiation codon, coding region, 5' or 3' end genemic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nuclectides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antinflammatory steroid and ubjusione. A composition of the invention has antinflammatory, antiallergic, antisthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a immunosuppressive, and cytostatic activity. The composition may have a creventing a respiratory, lung or malignant disease or condition, also preventing the prophylactic or therapeutic respiratory effect of an antisflammatory steroid in a subject, for reducing levels of denosine of or, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung surfactant in a subject's tissue, or treating bronchoconstriction, lung surfactant in a subject's tissue, or treating bronchoconstriction, lung surfactant in a subject's tissue, or treating bronchoconstriction, lung surfactant in a subject's tissue, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained pot_sequences

or fire, who only published pot_sequences 

Sequence 20 BP; 5 A; 6 C; 6 G; 3 T; 0 U; 0 Other;

Gaps ö / Match 3.1%; Score 13.2; DB 1; Length 20; Local Similarity 83.3%; Pred. No. 3.8e+02; nes 15; Conservative 0; Mismatches 3; Indels Query Match Best Loca Matches

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)369/c ABZ99369 standard; DNA; 20 BP. RESULT 392 ABZ99369/c

17-OCT-2003 (first entry) ABZ99369; 222222222 222222222

Human PDE4C oligonucleotide sequence.

Human, antisense, lung dysfunction, nasal airway dysfunction; antiinflammatory steroid; ubiquinone, antiinflammatory; antiallergic;

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antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens

WO200285308-A2

31-OCT-2002

23-APR-2002, 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Li Y, Sandrasagra A, Katz E, Pabalan J, Aguilar Tang L, Shahabuddin S; Nyce JW, I Miller S,

WPI; 2003-229219/22.

Pharmaceutical composition for treating allments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 14611; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation coodon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory teroid and ubiquinone. A composition of the invention has antiinflammatory steroid and ubiquinone. A composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an autistic ammatory steroid in a subject, for reducing levels of adenosine receptor, producing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject stissue, or treating bronchocomstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence date for this patent is not represented in the printed of specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences 

Sequence 20 BP; 1 A; 8 C; 8 G; 3 T; 0 U; 0 Other;

ö 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; ative 0; Mismatches 3; Indels 15; Conservative Local Similarity Query Match Best Loca Matches

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RESULT 393

ABZ92374 standard; DNA; 20

BP

ABZ92374;

(first entry) 17-0CT-2003

Human oligonucleotide sequence,

Human, antisense, lung dysfunction, nasal airway dysfunction, antiinflammatory steroid, ubiquinone, antiinflammatory, antiallergic, ABZ92374
ID ABZS
XX
XX
XX
DT 17-C
XX
XX
DDE Huma
XX
KW Huma
KW Anti

antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; broncholiation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Page 189

Homo sapiens

WO200285308-A2

31-0CT-2002

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC

Aguilar Li Y, Sandrasagra A, Katz E, Pabalan J, Tang L, Shahabuddin S; Nyce JW, I Miller S,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure, SEQ ID NO 7616, 872pp, English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligomucleotide antisense to the initiation codon, coding region, 5' or 3' or and genomic flanking regions; 5' and 3' intron-exon junctions, 5' or 3' or and genomic flanking regions; 5' and 3' intron-exon junctions, or regions within 2-10 mucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antistanatory, inng or malignant disease or condition, as so preventing a repliratory, inng or malignant disease or condition, as so in a reducing sensitivity to adenosine, reducing or depleting levels of adenosine receptor, producing bronchodialaion, increasing bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease; or condition or lung surfactant in a subject's tissue, or treating bronchoconstriction, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences 

Sequence 20 BP; 5 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

ô 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; ative 0; Mismatches 3; Indels Query Match 3.1 Best Local Similarity 83.3 Matches 15; Conservative

GAGGCTGGCCCTAAGAT 18 27 GAGGGCTGGGACGAGAT 44

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RESULT 394 ABZ97852 ID ABZ9785

踣. ABZ97852 standard; DNA; 20

ABZ97852;

(first entry) 17-0CT-2003

Human eotaxin oligonucleotide sequence.

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Human, antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic;

antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy, antisense gene therapy, respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds. 24-APR-2001; 2001US-0286137P. 23-APR-2002; 2002WO-US013135. EPIG-) EPIGENESIS PHARM INC. WO200285308-A2. Homo sapiens. 31-OCT-2002, 

Li Y, Sandrasagra A, Katz E, Pabalan J, Aguilar Tang L, Shahabuddin S; Nyce Jw, I Miller S,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 13094; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5, or 3 end genomic flanking regions, 5, and 3 intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubjunione. A composition of the invention has antiinflammatory, antiallargic, antiasthmatic, hypotensive, cue in antisense gene therapy. The composition may have a cue in antisense gene therapy. The composition may have a cue in antisense gene therapy. The composition may have a cue in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory stearid in a subject, for reducing or depleting levels of an entitial and subject in the subject, for reducing pronchodilation, increasing levels of ubjuninone or lung surfactant in a subject s tissue, or treating bronchoconstriction.

Colung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in alectronic format directly from WIPO cat fib., who, int/pub/published_pct_sequences

Sequence 20 BP; 1 A; 8 C; 4 G; 7 T; 0 U; 0 Other;

Gaps . Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels

308 CCCCGGGGACCGCGTGCT 325

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ABZ86905 standard; DNA; 20 ABZ86905; RESULT 399 ABZ86905/C XX ABZ8 AC ABZ8 DT 17-O XX Huma KW Huma KW Huma

BP.

(first entry) 17-OCT-2003

Human oligonucleotide seguence.

Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory; antiallergic;

antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

gene therapy;

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

Li Y, Sandrasagra A, Katz E, Pabalan J, Aguilar D; Tang L, Shahabuddin S; (EPIG-) EPIGENESIS PHARM INC Nyce JW,

WPI; 2003-229219/22.

Miller S,

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

claim 15; SEQ ID NO 2147; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the intitation coodon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of inactions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubjudinone. A composition of the invention has antiinflammatory antiallargic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic cativity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of according pronchodilation, increasing levels of adenosine ceptor, producing bronchodilation, increasing levels of adenosine creeptor, producing bronchodilation, increasing bronchoconstriction, lung surfactant in a subject s tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition.

Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO cet fits. 

Sequence 20 BP; 3 A; 3 C; 7 G; 7 T; 0 U; 0 Other;

0; Gaps 3.1%; Score 13.2; DB 1; Length 20; llarity 83.3%; Pred. No. 3.8e+02; Conservative 0; Mismatches 3; Indels Best Local Similarity Matches 15; Conserv Query Match

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RESULT 396 ACC47656

ACC47656 standard; DNA; 20

ACC47656;

16-SEP-2003 (first entry)

Human IGFBPS phosphorothioate antisense oligonucleotide, SEQ ID NO:32.

Human; insulin-like growth factor binding protein 5; IGFBP5; IBP5; chromosome 2q33-34; IGF signal transduction; IGF regulation; apoptosis; *****

/mcd_base= OTHER /note= "This oligonucleotide has a phosphorothicate /note= "This oligonucleotide (2'-MOE) wings at the 5' backbone and 2-'methyoxyethyl (2'-MOE) wings at the 5' and 3' ends, which are 5 nucleotides in length. Also all cytosine residues are 5-methylcytosines" bone growth stimulator; hyperproliferative disorder; cancer; tumour; breast; prostate; pancreas; neuroendocrine; inflammatory disorder; colitis; developmental disorder; growth disorder; buchenne muscular dystrophy; metabolic disorder; diabetes; osteoporosis; osteopetrosis; cytostatic; antiinflammatory; expression inhibition; phosphorothicate; antisense oligonucleotide; ss. Location/Qualifiers 1. .20 /*tag= a Key modified_base Homo sapiens

WO2003030826-A2

17-APR-2003

07-OCT-2002; 2002WO-US032060

09-OCT-2001; 2001US-00975123.

(ISIS-) ISIS PHARM INC.

Freier SM;

WPI; 2003-381673/36.

New antisense oligonucleotides for modulating insulin-like growth factor binding protein 5 gene expression, useful for preventing or treating cancers, inflammatory disorders, developmental disorders or metabolic cancers, ir disorders.

Claim 3; Page 76; 105pp; English.

Sequences ACC47637.ACC47667 represent phosphorothicate antisense cligonuclectides targeted to the human insulin-like growth factor binding protein 5 (IGFBPS) gane, which inhibit its expression. The antisense coligonuclectides were designed to target different regions of human coligonuclectides were designed to target different regions of human (IGFBPS RNA, and were analysed for their effect on IGFBPS) is a member of quantitative real-time PGR. IGFBPS (also known as IBPS) is a member of the insulin-like growth factor superfamily, which are involved in the crequiation of IGF action and bioavailabilty, and which also mediate IGFP independent actions, including inhibition or enhancement of apoptosis. IGFBPS is a key component of the IGF system in bone, having a high specific binding affinity for hydroxyapatite and extracellular matrix specific binding affinity for hydroxyapatite and extracellular matrix of specific binding affinity for hydroxyapatite and extracellular matrix of their an independent mechanism. IGFBPS is also expressed in other tissues, such as kidney, liver gut endothelium, lung tubules and other tissues, such as kidney, liver gut endothelium, lung tubules and other tissues, notechord, muscle cells in prostate cancer of colitis. It is also thought to play a role in prostate cancer of colitis. It is expressed with high frequency in neuroendocrine tumours, and has been shown to be induced in breast cancer cells upon treatment of progracledorides of the invention and treatment of information and treatment of information and treatment of information are useful for diagnosis, the propression is propressed in hyperproliferative disorders (particularly cancers of the breast, propertion and treatment of information are useful for diagnosis, or prosteders (e.g., colities) developmental or growth disorders (e.g., diabetes, colities, or osteoperosis)

Seguence 20 BP; 2 A; 8 C; 4 G; 6 T; 0 U; 0 Other;

Gaps . Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels

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CGAGGTCCTCAGTTTCCT 19 359 CGACTICCICACITICCI

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ACASS326 standard; DNA; 20 4CAS5326,

ACA55326;

05-JUN-2003 (first entry)

Human modified IgE CH2 domain PCR primer P171-A393

CH3 domain; IgE; antigen; non-anaphylactic; anti-IgE; fusion protein; dermatological; antiinflammatory; ophthalmological; allergy; asthma; allergy chinitis; gastroincestinal allergy; food allergy; eosinophilia; conjunctivitis; glomerular nephritis; flea allergy; atopic dermatitis; gene therapy; PCR; primer; se.

Homo sapiens. Synthetic.

EP1262491-A2.

04-DEC-2002.

22-MAY-2002; 2002EP-00253606.

22-MAY-2001; 2001US-0292638P.

(PFIZ ) PFIZER PROD INC.

Brown TM, Morsey MA;

WPI; 2003-122561/12.

Novel isolated antigenic peptide comprising amino acid residues of CH3 domain of IgE molecule from first species and a second unrelated species, induces non-anaphylactic anti-IgE immune response in animal.

Example 2; Page 22; 50pp; English.

This invention describes a novel antigenic peptide comprising amino acid residues of an IgB CH3 domain from a first species (ADE1) and amino acid residues of an IgB CH3 domain from a first species (ADE2), where ADE1 is conserved in the IgE CH3 domain of the second apecies and ADE2 is not conserved in the IgE CH3 domain of the second apecies and ADE2 is antigenic peptide induces a non-anaphylactic anti-IgE immune response in a national. The invention also discloses the polynucleotide sequence conciding the antigenic peptide and an antigenic fusion protein comprising the antigenic peptide of the invention and a heterologous protein comprising the antigenic peptide of the invention and a heterologous protein comprision carrier, where the fusion protein induces an anti-IgE immune response that does not cause anaphylaxis when administered to an animal. The corputors of the invention have dermatological, antiinflammatory and ophthalmological activity. The antigenic peptide described is capable of products of the invention are useful in the manufacture of a medicament for treating or preventing IgE-mediated allergic disorders including asthma, allergic rhinitis, gastrointestinal allergic disorders including asthma, allergic rhinitis, gastrointestinal allergic disorders consoluted products are useful for treating IgE-mediated allergic disorders by gene therapy. Antigenic peptides comprising conserved amino conditive by variable amino acid residues of the CH3 domain of an IgE molecule from ans species cald residues of fanti-IgE antibodies when administered to an animal without causing anaphylaxis. ACAS5184-ACAS5346 represent PCR primers used to amplify the polymucleotide sequences used in designing the constructs causing anaphylaxis. ACAS5184-PEPESET Conserved in the disclosure of the invention 

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The invention relates to a compound 8-50 nucleobases in length targeted to a nucleic acid molecule encoding dual specific phosphatese 5, where the compound specifically hybridises with and inhibits the expression of dual specific phosphatese 5. The compound is used for treating an animal having a disease or condition associated with dual specific phosphatese 5 such as a hyperproliferative disorder, a developmental disorder, an inflammatory disorder or a disease which arises from aberrant apoptosis. Sequences ARX09062-ABX09139 represent human dual specific phosphatese 5 phosphorothioate oligonucleotides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cytostatic; immunostimulant; 162P1E6; cytotoxic agent; immune response; cancer; bladder; prostate; kidney; lung; breast; passive immunisation; transgenic animal; vaccine; gene therapy; PCR; primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antisense modulation of dual specific phosphatase 5 expression used in treating disorders e.g. inflammatory diseases.
                                                                                                                                                                             Human; dual specific phosphatase 5; ss; developmental disorder; hyperproliferative disorder; inflammatory disorder aberrant apoptosis; antiinflammatory; cytostatic; antiapoptotic; antiproliferative; phosphorothicate oligonucleotide.
                                                                                                                                           Human dual specific phosphatase 5 phosphorothioate oligonucleotide #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 2 A; 9 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    162P1E6 cancer gene related nested primer NP2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 15; Page 84; 110pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       387 GACGCCCCAAGAAGGTC 404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20 GGCGGCGCATGAAGGTC 3
         063/c
ABX09063 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-MAY-2002; 2002WO-US015305
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Best Local Similarity 83.3
                                                                                                        22-JAN-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ISIS-) ISIS PHARM INC
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                                                                                                                                                                                                                                                                                                      Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABT17425;
                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                  ABX09063;
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ABT17425/c
셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a novel composition comprising a substance that modulates the status of a 1519304 protein (e.g. 1519304 variant 1-11; or a molecule that is modulated by the 1519304 protein, where the status of compositions, or the 1519304 protein is modulated. The novel compositions, or the 1519304 proteins and genes, are useful for eliciting a humoural or cellular immune response. The 1519304 genes and proteins are also useful for diagnosing, prognosing, preventing or treating cancer, e.g. adenocarcinoma, bladdar cancer, colorectal cancer, lung or bronchial cancer, breast cancer or carcinoma. This polymucleotide sequence represents a 1519304 related primer of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New 151P3D4 proteins and genes, useful for eliciting a humoral or cellular immune response, or for diagnosing, prognosing, preventing or treating cancer, e.g. adenocarcinoma, bladder cancer, lung, breast cancer
                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cytostatic; gene therapy; vaccine; modulator; 15193D4; humoural; cancer; cellular immune response; adenocarcinoma; bladder; colorectal; lung; bronchial; breast; carcinoma; PCR; primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                           Gaps
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                                             3.1%; Score 13.2; DB 1; Length 20;
83.3%; Pred. No. 3.8e+02;
Live 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
      Sequence 20 BP; 2 A; 6 C; 4 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 1; Page 69; 426pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Challita-Eid PM, Raitano AB, Far
Morrison RK, Ge W, Jakobovits A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         373 TCCTGGACCGCGACG 390
                                                                                                                                    384 GACGACGCCCAAGAAG 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TCCTCGGCCGCGACCACG 3
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                                                                                                                                                                                                                                                                                      ABT43860 standard; DNA; 20 BP.
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25-APR-2001; 2001US-0286630P.
                                                                                                                                                                                                                                                                                                                                                                                                                   DPNCDN nested primer 2 (NP2)
                                                                                                                                                                        GAGGACACCAAGAAG
                                                                                                                                                                                                                                                                                                                                                                        16-OCT-2003 (first entry)
                                               Query Match
Best Local Similarity 83.33
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (AGEN-) AGENSYS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-167091/16.
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Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200283860-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     or carcinoma.
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Unidentified.

ABT43860;

ABT43860/

13

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Gaps ö

20

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RESULT 399

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The invention relates to a novel composition comprising a substance that modulates the status of a 162P1E6 protein. The protein comprises one of 21 sequences of 70 - 146 amino acids, given in the specification, or a molecule that is modulated by the protein, where the status of a cell that expresses the protein is modulated. An antibody to the 162P1E6 protein is used to deliver a cytotoxic agent or a diagnostic agent to a cell that expresses the 162P1E6 protein. The composition is used for centering the presence of a 162P1E6 related protein or a 162P1E6 related protein as 162P1E6 related proteins and polymuclectide in a sample. The 162P1E6 proteins and polymuclectide in a sample. The 162P1E6 proteins and polymuclectide senceding them are useful for diagnosing, proteins and polymuclectides encoding them are useful for adaptosing, prostate cancer, kidney cancer, lung cancer, or breast cancer. They can also be used for eliciting a humoral or cellular immune response. The mutibodies or T cells reactive with 162P1E6 are useful for active or passive immunisation. Transgenic animals are useful for developing and screening of useful reagents. The polymuclectide and polympetide cancer of the invention can also be used to treat disorders by being used in a vaccine or in gene therapy. This polymuclectide sequence
                                                                                                                                                                                                                                                                                                                                                              Composition for diagnosing, prognosing, preventing or treating cancer, for eliciting a humoral or cellular immune response, or for active or passive immunization, comprises a substance that modulates the status of
                                                                                                                                                                                                                                                   Challita-Eid PM, Raitano AB, Faris M, Hubert RS, Morrison K;
Morrison RK, Ge W, Jakobovits A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 71; 437pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       373 TCCTGGACCGCGACG 390
                                                                                       09-APR-2002; 2002WO-US011544.
                                                                                                                                 10-APR-2001; 2001US-0283112P
25-APR-2001; 2001US-0286630P
                                                                                                                                                                                                                                                                                                                       WPI; 2003-148268/14.
                                                                                                                                                                                                          (AGEN-) AGENSYS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                     a 162P1E6 protein.
WO200283916-A2
                                              24-OCT-2002
à
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0; Gaps
3.1%; Score 13.2; DB 1; Length 20; ilarity 83.3%; Pred. No. 3.8e+02; Conservative 0; Mismatches 3; Indels
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20 recresececesaceace 3 셤

.621/c ACD02621 standard; DNA; 20 BP. 31-JUL-2003 (first entry) ACD02621; 

Suppressive subtractive hybridisation of STEAP related primer #8.

STEAP-1; six transmembrane epithelial antigen of the prostate; cancer; cancer vaccine; delineation; cytogenetic abnormality; cytostatic; vaccine; PCR; primer; ss.

Homo sapiens

WO2003022995-A2.

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The invention describes a composition comprising a substance that modulates the status of a protein (1) of 340 or 283 amino acids, or of any of the 15 sequences of 259 amino acids, given in the specification, or a molecule that is modulated by the protein, where the status of the cell that expresses the protein is modulated. The compositions, proteins, polynucleotides and methods are useful for treating and detecting cancer. The STEAP-1-related proteins are useful for generating cancer vaccines. The polynucleotides are useful as tools for delineating, with greater precision, cytogenetic abnormalities in the chromosomal region that encodes STEAP-1 that may contribute to the malignant phenotype. This sequence represents a primer used to analyse human six transmembrane epithelial antigen of the prostate or STEAP-1 cDNA's
                                                                                                                                                                                                                                                                 New composition comprising a substance that modulates the status of STEAP-1-related protein, useful for treating and detecting cancer.
                                                                                                                                                                                      Jakobovits A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                      Challita-Eid PM,
                                                                                                                                                                                                                                                                                                                                  Example 1; Page 70; 248pp; English.
                                                                                                                                                                                      Raitano AB,
                                        06-SEP-2002; 2002WO-US028371.
                                                                                06-SEP-2001; 2001US-0317840P.
05-APR-2002; 2002US-0370387P.
                                                                                                                                                                                                                            WPI; 2003-313240/30.
                                                                                                                                              (AGEN-) AGENSYS INC.
                                                                                                                                                                                      Ge ₩,
20-MAR-2003
                                                                                                                                                                                      Faris M,
```

Gaps ö Score 13.2; DB 1; Length 20; Pred. No. 3.8e+02; 0; Mismatches 3; Indels 3; Indels ö 373 TCCTGGACCGCGACG 390 20 TCTCGGCGCGACCACG 3 3.1%; Query Match
Best Local Similarity 83.3
Matches 15, Conservative g ò

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88 Cytostatic; vaccine; cancer; immune response; PCR; primer; 

Hubert RS; Ge ≅ A, Challita-Eid PM, Faris M, Morrison RK, Raitano AB;

New composition comprising a substance that modulates the structure of proteins and polynucleotides, useful for therapeutic, prognostic and diagnostic reagents for eliciting cellular or humoral immune response in cancer patients.

Example 1; Page 72; 1021pp; English.

The present invention relates to novel human cancer-related genes and proteins (ABZ78120-ABZ78168 and ABR01789-ABR01861). The genes and proteins are useful for eliciting a humoral or cellular immune response. The genes are useful as probes and primers for the amplification and/or detection of genes, mRNAs or their fragments, as reagents for the detection of genes, mRNAs or their fragments, as reagents for the diagnosis and/or prognosis of cancer, as coding sequences capable of directing the expression of the protein, as tools for modulating or inhibiting the expression of genes and/or translation of transcripts, and as therapeutic agents. The proteins and peptides are useful as therapeutic, prognostic and diagnostic reagents for cancer. The present sequence is a primer, used in an example from the invention

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

. 0 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.88+02; ative 0; Mismatches 3; Indels 373 TCCTGGACCGCGACGACG 390 20 TCCTCGGCGCGACCACG 3 Local Similarity 83.3 Query Match Matches ઠે

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Gaps

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Cancer associated coding sequence PCR primer #3.

Cancer associated coding sequence; cancer; human; cytostatic; gene therapy; PCR; primer; ss.

10-APR-2001; 2001US-0282739P. 10-APR-2001; 2001US-0283112P. 25-APR-2001; 2001US-028630P. 10-APR-2002; 2002US-00286630P.

10-APR-2002; 2002WO-US011645.

Challita-Eid

New pharmaceutical composition for diagnosing, prognosing, preventing or treating cancer, comprises a substance that modulates a nucleic acid sequence, e.g. 105P1B7, 152P1A2B or 156P3A6, or a molecule modulated by the nucleic acid.

ğ

The invention comprises the amino acid and coding sequence of a 184PIE2 protein. The DNA and protein sequences of the invention are useful for diagnosing, prognosing and/or treating cancer. The 184PIE2 DNA and protein sequences may also be used to elicit a humoral or a cellular immune response in patients and in monitoring genetic abnormalities. Antibodies raised against the 184PIE2 proteins may be use in active or passive immunisation. The present DNA sequence is used in the exemplification of the invention

Example 1, Page 69; 394pp; English.

Example 1, Page 34; 72pp; English.

The present invention relates to a pharmaceutical composition comprising a substance that modulates the status of a cancer associated mucleic acid sequence such as given in the specification (see ABZ20564-ABZ20575) or a molecule that is modulated by the above nucleic acid sequence, where the 

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Gaps

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Query Match 3.1%; Score 13.2; DB 1; Length Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

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status of a cell that expresses the nucleic acid sequence is modulated. The composition is useful in diagnosing, prognosing, preventing and/or treating ancer. The nucleic acid sequence may be used in monitoring genetic abnormalities, in generating and characteriaing domain-specific antibodies, for identifying agents or cellular factors that bind to a protein, and in therapeutic and diagnostic contexts, such as diagnostic seasys, cancer vaccines, and methods of preparing vaccines. The present sequence is a primer used to identify the cancer associated coding sequences suitable to be modulated in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    useful for
in eliciting an
                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Faris M, Hubert RS, Morrison
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gene therapy, vaccine, 184P1E2, cancer, genetic abnormality; cellular immune response; immunisation; PCR; primer; ss.
                                                                                                                                                                           Score 13.2; DB 1; Length 20;
Pred. No. 3.8e+02;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New 184PIE2 polynucleotide encoding a 184PIE2 protein, diagnosing, prognosing, preventing or treating cancer, immune response, and in chromosome mapping.
                                                                                                                                                   Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                               184P1E2 gene-specific nested PCR primer #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chalitta-Eid PM, Raitano AB, Fari
Morrison RK, Ge W, Jakobovits A;
                                                                                                                                                                                                                                       373 TCCTGGACCGCGACG 390
                                                                                                                                                                                                                                                                  m
                                                                                                                                                                                                                                                                                                                                           BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-APR-2001; 2001US-0282739P.
25-APR-2001; 2001US-0286630P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-APR-2002; 2002WO-US011643
                                                                                                                                                                               Query Match 3.1%;
Best Local Similarity 83.3%;
Matches 15; Conservative
                                                                                                                                                                                                                                                            20 TCCTCGGCCGCGACCACG
                                                                                                                                                                                                                                                                                                                                         AAL52254 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-148269/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (AGEN-) AGENSYS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200283919-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                   16-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24-OCT-2002.
                                                                                                                                                                                                                                                                                                                                                                         AAL52254;
                                                                                                                                                                                                                                                                                                             RESULT 40
AAL52254/
                                                                                                                                                                                                                                                                                                                                              888888888888888
                                                                                                                                                                                                                                                                  셤
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Length 20;

Query Match

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The invention relates to antisense compounds targetted to a nucleic acid molecule encoding fibroblast growth factor (FGF) receptor 3 (also known as FGFR-3) ACH, UTK4 and CEK2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with FGFR-3 such as developmental disorders or hyperproliferative disorders, especially cancer of colorateds. bone long, cervical, breast or skin. They are useful as research reagents, therapeutics, prophylaxis, kits and diagnostics, and as tools in differential and/or combinatorial analyses to elucidate expression patterns of a portion of the genes expressed within cells and tissues. They are also useful in antisense therapy. The present sequence is an antisense oligonucleotide targetted to human FGFR-3
                                                                                                                                                                                                                                                                                                                                                             1..20
/*tag= a
//note= "Phosphorothioate backbone; All cytidine residues
are 5-methylcytidines"
                                                                                                                                                                                                                                Human, antisense; fibroblast growth factor receptor 3; prophylaxis;
developmental disorder; hyperproliferative disorder; antisense therapy;
FGFR-3; ACH; JTK4; CEK2; cancer; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel compound targeted to a nucleic acid molecule encoding fibroblast growth factor receptor 3, useful for inhibiting the expression of the receptor and for treating an animal having cancer or developmental
                                                                                                                                                                                                                                                                                                                                                                                                                                                     /*tag= b
/mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
15. 20
/*tag= c
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /mod_base= OTHER
/note= "2 -methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                     Human FGFR-3 antisense oligonucleotide, ISIS #125169.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 15; Page 79; 120pp; English
                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
373 TCCTGGACCGCGACGACG 390
                465/c
AAD55465 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-SEP-2002; 2002WO-US028549.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-SEP-2001; 2001US-00953047
                                                                                                                                                                      07-AUG-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Monia BP, Wyatt JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-313244/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO2003023004-A2
                                                                                                                                                                                                                                                                                                                                              Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                         modified base
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                                                                                                                                                                                                                                                                                                                  Synthetic
                             20
                                                                                                                                        AAD55465;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     disorder
                                                                                         AAD55465,
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Seguence 20 BP; 5 A; 6 C; 7 G; 2 T; 0 U; 0 Other;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes a compound (1) 8-50 nucleobases in length targeted to a nucleic acid molecule encoding BCL2-associated X (BAX) procein, where the compound specifically hybridises with the nucleic acid molecule encoding BAX protein and inhibits the expression of BAX protein. The compound specifically hybridises with at least 8-nucleobase portion of an active site on a nucleic acid molecule encoding BAX protein. Also described: (1) a composition comprising (1) and a pharmaceutical carrier
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New antisense compounds, useful for modulating the expression of BCL2-associated X (BAX) protein or for treating a disease or condition associated with BAX protein, e.g. Parkinson's disease, Hodgkin's disease or Alzheimer's disease.
                                                                                                                                                                                                                                                                               BCL2-associated X; BAX; nootropic; neuroprotective; antiparkinsonian; anticonvulsant; ophthalmological; antidiabetic; virucide; antisense therapy; BAX antagonist; BAX inhibitor; familial amylotrophic lateral sclerosis; Alzheimer's disease; Parkinson's disease; Hodgin's disease; satillage-hair hyperplasia; diabetes-associated ocular disorder; scrapie infection; aberrant apoptosis; human; phosphorothioate; ss.
                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mcd_base= OTHER
/note= "phosphorothioate linkages, and all cytidine
residues are 5-methylcytidines"
                                                                                                                                                                                                                                                    Human BAX chimeric phosphorothioate oligonucleotide SEQ ID NO:26
                             ö
                             3; Indela
Score 13.2; DB 1;
Pred. No. 3.8e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /*tag= a
/mod_base= OTHER
/note= "2'-0-methoxyethyls"
16.20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /*tag= c
/mod_base= OTHER
/note= "2'-0-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 3; Page 85; 139pp; English.
                                                          54 TCAGAGGAGTCTCTGCAC 71
                                                                                   19 TCAGAGGGCCTCTGCTC 2
                                                                                                                                                                ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13-JUL-2002; 2002WO-US022417.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-JUL-2001; 2001US-00908147.
3.1%;
                                                                                                                                                                ADA20853 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Д
                                                                                                                                                                                                                        20-NOV-2003 (first entry)
                               15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-239321/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Zhang H, Watt AT;
                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO2003008543-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Key
modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-JAN-2003.
                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                             ADA20853;
                  Best Local
Matches 1
                                                                                                                                   RESULT 406
                                                                                                                                                 ADA20853
ID ADA2
                                                                                                                                                                               셤
                                                             ઠે
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tissues comprising contacting the expression of BAX protein in cells or tissues comprising contacting the cells or tissues with (I); and (3) creating an animal having a disease or condition associated with BAX protein comprising administering to the animal (I) so that expression of BAX protein is inhibited. (I) has nootropic, neuroprotective, antiparkinsonian, anticonvulsant, ophthalmological, antidabetic and virudde activities, and can be used in antissense therapy, and as a BAX cragonist. The antissense compounds (I) are useful for modulating the expression of BAX protein, and for treating a disease or condition associated with BAX protein, e.g. familial amylotrophic lateral cartiage-hair hyperplasia, disbetes-associated ocular disorders or scrapie infection, or a condition that arises from abbrrant apoptosis. The compounds are useful as research reagents and in diagnostics. The present sequence represents a human BAX chimeric phosphorothioate oligonucleotide, which is used in an example from the present invention.

Sequence 20 BP; 1 A; 9 C; 7 G; 3 T; 0 U; 0 Other;

3; Indels Ouery Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels ò ద

ADA26274 standard; DNA; 20 BP RESULT 407 ADA26274/c

ADA26274;

Zebrafish genomic DNA PCR primer #2.

20-NOV-2003 (first entry)

Zebrafish, PCR; ss; hedgehog; neuronal cell; skeletogenesis; chrondrogenesis; osteogenesis; degenerative disorder; nervous system; neuronal cell death; neural cell; neuromuscular disorder; autonomic disorder; central nervous system disorder; anoxia; ischaemia; peripheral nervous system disorder; tachycardia; atrial cardiac arrhythmia; striated heart; stem cell development; digestive tract; liver; multiple sclerosis; primer. 

Danio rerio.

US2003054437-A1.

20-MAR-2003

97US-00954771.

20-OCT-1997;

93US-00176427. 94US-00356060. 95US-00435093. 95US-00462386. 30-DEC-1993; 14-DEC-1994; 04-MAY-1995; 05-JUN-1995;

(INGH/) INGHAM P W. (MCMA/) MCMAHON A P. (TABI/) TABIN C J.

Tabin CJ; ingham PW, Mcmahon AP,

WPI; 2003-555377/52

Modulating growth, differentiation or survival of a cell, useful for treating a degenerative disorder of the nervous system characterized by neuronal cell death, comprises contacting the cell with a hedgehog polypeptide.

Example 4; Page 44; 121pp; English

The invention relates to a method for modulating growth, differentiation or survival of a cell, comprising contacting the cell with a hedgehog polypeptide. The invention also relates to methods for inducing a cell to differentiate to a neuronal cell phenotype comprising a cell to differentiate to a neuronal cell phenotype comprising contacting a cell to carget tissue of a hedgehog polypeptide comprising a degenerative disorder of the nervous system characterised by neuronal cell death, comprising administering a hedgehog polypeptide causing prolonged survival of neural cells in the patient, relative to the absence of hedgehog treatment. The hedgehog polypeptide causing prolonged survival of neural cells in the patient, relative to the absence of hedgehog treatment. The hedgehog polypeptides are useful for treating a degenerative disorder of the nervous system characterised by neuronal cell death, including contents a clerosis, pick's disease, Parkinson's disease, amyotrophic specifically Alzheimer's disease, Parkinson's disease, multiple contents and correction associated with a natural aging process. The neuronal degeneration associated with a natural aging process. The neuronal degeneration associated with a natural aging process. The collisor including disorders affecting innervation of smooth muscle and disorders affecting innervation of smooth muscle and peripheral nerve damage, for treating neoplastic or hyperplastic carriythmias which and peripheral nerve damage, for treating neoplastic or hyperplastic carriythmias which and peripheral nerve damage, for treating neoplastic or hyperplastic carriythmia such a degeneration as disease. See the cells of the heart, in nerve proscheses for repairing central cargine from a degeneration of the digestive tract, liver and other organs. This sequence represents a PCR primer used to amplify zebrafish %\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$

Sequence 20 BP; 3 A; 6 C; 2 G; 1 T; 0 U; 8 Other;

Gaps ., 3.1%; Score 13.2; DB 1; Length 20; 60.0%; Pred. No. 3.8e+02; rative 2; Mismatches 6; Indels Best Local Similarity 60.0 Matches 12, Conservative Query Match

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ઠ 쉺 RESULT 408 ACF57119

BP. ACF57119 standard; DNA; 20 ACF57119; Human sulfatase related probe SEQ ID NO:9.

14-OCT-2003 (first entry)

Human; sulfatase; enzyme; cytostatic; neuroprotective; nootropic; antiparkinsonian; cerebroprotective; analgesic; cardiovascular; cardiant; antiansonian; antianthumic; antiantemic; nephrotropic; uropathic; antiinflammatory; vasotropic; antiathmatic; gene therapy; cancer; CNS disorder; COPD; central nervous system disorder; cardiovascular disorder; asthma; haematological disorder; genitourinary disorder; chromosome X; Xp22.33; chronic obstructive pulmonary disease; probe; ss. PART SERVICE S

Homo sapiens. Synthetic.

WO2003057869-A1.

17-JUL-2003.

09-JAN-2003; 2003WO-EP000137

14-JAN-2002; 2002US-0347247P. 29-JUL-2002; 2002US-0398732P.

(FARB ) BAYER AG

Liou J;

rng.res

The present invention describes a human sulfatase enzyme (I), which is located on chromosome X (more specifically to Xp22.33). (I) has cardiant, concreptor, cytostatic, neuroprotective, antiparkinsonian, analgesic, cerebroprotective, cardiovascular, antiarrhythmic, antianeamic, cerebroprotective, cardiovascular, antiarrhythmic, antianeamic, activities, and can be used in gene therapy. The sulfatase polymucleotide and polypeptide sequences can be used in diagnosing, preventing, ameliorating or treating diseases associated with sulfatase dysfunction. They may also be used to identify test compounds that may act, for example, as activators or inhibitors at the enzyme's active site. The human sulfatase and its fragments are also useful in raising specific antibodies that can block the enzyme and effectively reduce its activity. The sulfatase can be used as hybridiation probes or primers. The sulfatase can be used as hybridiation probes or primers. The sulfatase can be used as hybridiation probes or primers. The sulfatase can be used in the treatment of diseases such as cancer, a central nervous system (CNS) disorder (e.g. Alzheimer's disease, concer, a disorder (e.g. heart failure or arrhythmias), a haematological disorder (e.g. anaemia or thrombocytopaenia), a genitourinary disorder (e.g. renal failure) obstructive pulmonary disease (COPD) or asthma. The present sequence represents a probe for human sulfatase, which is used in an example from the present invention New polynucleotide encoding a sulfatase polypeptide, useful for diagnosing, preventing or treating diseases associated with sulfatase dysfunction, e.g. cancer, asthma or cardiovascular disorders. Example 16; Page 99; 124pp; English

Sequence 20 BP; 4 A; 8 C; 5 G; 3 T; 0 U; 0 Other;

Score 13.2; DB 1; Length 20; Pred. No. 3.8e+02; 0; Mismatches 3; Indels ò 3.1%; Query Match
Best Local Similarity 83.3
Matches 15; Conservative ઠે 셤

ACD44752 standard; DNA; 20 BP 09-SEP-2003 ACD44752; RESULT 40 

(first entry)

PKA regulatory subunit RII alpha inhibitory oligonucleotide ISIS102778.

Human; ss; antisense therapy; infection; inflammation; tumour; protein kinase A regulatory subunit RII alpha.

Synthetic. Homo sapiens.

US6524854-B1

25-FEB-2003

11-SEP-2001; 2001US-00954560

11-SEP-2001; 2001US-00954560.

(ISIS-) ISIS PHARM INC

Cowsert Monia BP,

WPI; 2003-511923/48.

New antisense compounds, useful for modulating the expression of protein kinase A (PKA) regulatory subunit RII alpha, and for treating a disease or condition associated with expression of PKA regulatory subunit RII alpha. ####X%X00000000000X%

Example 15; Col 43-44; 35pp; English

The invention relates to antisense compounds targeted to nucleic acids encoding protein kinase A regulatory subunit RII alpha. The antisense compounds are useful for modulating the expression of protein kinase A (PKA) regulatory subunit RII alpha and for treating a disease or condition associated with expression of PKA regulatory subunit RII alpha. The compounds are also useful as research reagents and kits, or for diagnostics, therapeutics and prophylaxis, e.g. to prevent or delay infection, inflammation or tumour formation. The present sequence represents a human protein kinase A regulatory subunit RII alpha inhibitory oligomucleotide

Sequence 20 BP; 3 A; 7 C; 9 G; 1 T; 0 U; 0 Other;

Gaps . 0 Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels

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320 CGTGCTGGCGGCGACGA 337

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ADB89866 standard; DNA; 20 RESULT 410 ADB89866/c ID ADB898

BP.

ADB89866;

04-DEC-2003 (first entry)

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Gaps ; 0

Antinsense oligonucleotide targeting human C3 component, ISIS139968.

Human; ss; antisense; complement component C3; inflammation; septic shock; multiple organ failure; hyperacute organ failure; autoimmune disorder; CNS inflammation; multiple sclerosis; tatherosclerosis; tumour.

Homo sapiens

Location/Qualifiers Ω Key modified base 

/mod_base= OTHER /note= "Phosphorothioate backbone and all cytosines are 5 -methyl cytosines" modified_base

/*tag* a /mod base= OTHER /note= "2'-methoxyethyl nucleotides" 16. .20 modified_base

*tag= c /mod_base= OTHER /note= "2'-methoxyethyl nucleotides"

US2003096775-A1

22-MAY-2003.

23-OCT-2001; 2001US-00001076

23-OCT-2001; 2001US-00001076.

SISI (-SISI)

Graham MJ,

rng.res

WPI; 2003-606441/57. 

```
New antisense oligonucleotides targeted to a nucleic acid molecule encoding complement component C3, useful for treating a disease or condition associated with complement component C3, e.g. autoimmune
                                                                                                                                                                  disorder or infection.
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Claim 3; Page 25; 72pp; English

The invention relates to a compound 8-50 nucleobases in length targeted to a nucleic acid molecule encoding complement component C3. The compound specifically hybridises with the nucleic acid molecule encoding complement component C3, or specifically hybridises with at least an 8-nucleobase component C3. Also included are a composition of complement component C3. Also included are a composition comprising the component C3 in cells or tissues (comprising the expression of complement component C3 in cells or tissues (comprising complement C3 in cells or tissues) component C3 in cells or tissues of complement component C3 in cells or tissues of complement component C3 compound cited above) and treating an animal cappound administering to the animal the compound cited above so that expression of complement component C3 is inhibited. The antisense compounds are useful for inhibiting the expression of complement C3 in cells or tissues, or for treating an animal having a cutoimmune disorder (e.g. multiple solerosis), an infection, or atherosclerosis, inflammation, septic shock, multiple organ failure, component C3 in distinguishing functions of various members of a biological pathway, or for preventing or delaying confidence or infection, inflammation or tumour formation. The present sequence is an infection, inflammation or tumour formation. The present sequence is an inferior. antisense oligonucleotide targeting human C3

Sequence 20 BP; 3 A; 8 C; 6 G; 3 T; 0 U; 0 Other;

ö Gaps ö ch 1 Similarity 83.3%; Pred. No. 3.8e+02; 15; Conservative 0; Mismatches 3; Indels Query Match Best Local Similarity Matches

321 GTGCTGGCGGCGGACGAC 338

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18 GIGCIGGAGGCCCACGAC 1

ADB68562 standard; DNA; 20 BP. RESULT 41 ADB68562/ 

ADB68562;

(first entry) 04-DEC-2003 homogeneous conjugate; hepatic; chronic viral hepatitis; cirrhosis; malaria; viral infection; protozoan; cancer; hepatocellular carcinoma; HCC, HCV; SB.

DNA oligonucleotide 9 targeted to Hepatitis C virus sequence.

Hepatitis C virus.

402003067209-A2

14-AUG-2003.

21-JUN-2002; 2002WO-US019908

2-JUN-2001; 2001US-00888164.

(CELL-) CELL WORKS INC. (UYJO ) UNIV JOHNS HOPKINS.

ΰ ຜ Deamond Duff R, Zhou Y, rs'o POP,

WPI; 2003-697456/66

New homogeneous prodrug conjugate containing hepatic ligand for delivery of pathogen-specific oligomer useful for treating liver infections or

Disclosure, Page 23; 107pp; English

The invention relates to a novel homogeneous conjugate comprising a hepatic ligand, bifunctional linker and biologically stable oligomer that binds to a sequence in a hepatic virus or pathogen and is released from the conjugate by hydrolysis or reduction. The conjugate of the invention may be useful during the treatment of liver diseases including chronic viral hepatitis, cirrhosis, malaria, viral or protozoan infection and cancer, such as hepatocellular carcinoma (HCC). The current sequence is that of the DNA oligomucleotide 9 of the invention which is targeted to a Hepatitis C virus (HCV) sequence. 

Sequence 20 BP; 2 A; 6 C; 6 G; 6 T; 0 U; 0 Other;

ö 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; ative 0; Mismatches 3; Indels Ouery Match
Best Local Similarity 83.5.
Best Local Similarity
The Conservative

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RESULT 412 ADC71183/c

ADC71183 standard; DNA; 20 BP.

ADC71183;

(first entry) 18-DEC-2003 Nested PCR primer 2 (NP2) used in SSH to isolate 205P1B5 cDNA fragment.

205P1B5; prostate cancer; immune response; transgenic; knock out animal; cytostatic; immunogenic; vaccine; ss; SSH; suppressive subtractive hybridisation; PCR; primer; NP2.

Unidentified

WO2003020954-A2

13-MAR-2003.

30-AUG-2002; 2002WO-US027760.

31-AUG-2001; 2001US-0316664P.

(AGEN-) AGENSYS INC.

Jakobovits A; Hubert RS, Faris M, Raitano AB, Challita-Eid PM, 

WPI; 2003-354484/33.

New polynucleotide designated 205PlBS, for diagnosing and treating prostate cancer, and as probes or primers for the amplification and/or detection of 205PlBS genes.

Example 1; Page 60; 162pp; English.

This invention relates to a novel gene designated 205P1B5, and the encoded protein, which is aberrantly expressed in prostate cancer. Specifically, it refers to the two variants of 205P1B5 mapped to chromosome 8p21-8p12, namely 205P1B5v1 and 205P1B5v2 and fragments thereof that serve as useful diagnostic, prophylactic, prognostic, prognostic and/ or therapeutic targets for prostate and other types of cancers. The present invention describes methods for the isolation of 205P1B5, for generating an immune response and for generating transgenic or knock out animals for

Sequence 20 BP; 3 A; 3 C; 7 G; 7 T; 0 U; 0 Other;

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the development and screening of therapeutically useful reagents. Furthermore, it refers to identifying proteins, small molecules or other agents that interact with 205P185, and can be used to identify pathways activated by 205P185. Accordingly, these are cytostatic and immunogenic compositions that are useful for the development of cancer vaccines. This suppressive subtractive is the nested PCR primer 2 (NP2) used for suppressive subtractive hybridisation (SSH) to isolate the 205P185 cDNA fragment of the invention.

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

Gaps ö 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; rative 0; Mismatches 3; Indels 373 TCCTGGACCGCGACGACG 390 m 20 rccrccccccccacc Query Match Best Local Similarity 83.3³ Matches 15, Conservative

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ADC16779 standard; DNA; 20 BP ADC16779; RESULT 413 ADC16779/ 

18-DEC-2003 (first entry)

Forward RT-PCR primer to amplify HIV-1 RNA in order clone T-20.

RT-PCR; PCR; primer; anti-retroviral; T-20; T-1249; 5-Helix; env; gp41; anti-HIV; vaccine; albumin fusion protein; HIV fusion inhibiting peptide; ss; cyanovirin-N

Unidentified

WO2003066078-A1.

14-AUG-2003.

07-FEB-2003; 2003WO-IB000434.

07-FEB-2002; 2002US-0355547P.

(AVET ) AVENTIS BEHRING GMBH. (DELZ ) DELTA BIOTECHNOLOGY LTD.

Hauser H, Weimer T, Sleep D;

WPI; 2003-731478/69.

New albumin fusion protein comprising a human immunodeficiency virus (HIV) fusion inhibiting peptide and an albumin having an albumin activity, useful for treating a disease or disorder, e.g. HIV infection.

Example 3; Page 58; 105pp; English.

This invention relates to novel albumin fusion proteins comprising a human immunodeficiency virus (HIV) fusion inhibiting peptide, which exhibit anti-retroviral activity. Specifically, it refers to inhibitory peptides including T-20, T-1249, 5-Helix or cyanovirin-N that bind the HIV env protein, or derivatives thereof such as the HIV gp41 protein. Furthermore, the albumin activity has the ability to prolong the in vivo half-life of these HIV fusion inhibiting peptides. Accordingly, the present invention describes fusion proteins that neutralise HIV in a host of prince in mimune response and also antibodies that inhibit viral infection of uninfected cells. In this way, a method exists to prevent, treat or ameliorate HIV infection and/ or a disease caused by HIV infection and/ or a disease caused by HIV infection and/ or a disease caused by HIV infection and or a disease caused by HIV activity and can be used towards the production of a vaccine. This oligonucleotide sequence is the forward RI-PCR primer used to amplify the HIV-1 RNA in order to clone T-20, in an exemplification of the invention.

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                                                                                                                       cytostatic; gynaecological; endometriosis; endometriotic haptoglobin;
ENDO-1; rat; PCR; primer; ss.
             Gaps
             ;
0
Length 20;
             Indels
                                                                                                             Rat endometriotic haptoglobin ENDO-1 primer seq id 7.
Match 3.1%; Score 13.2; DB 1; Local Similarity 83.3%; Pred. No. 3.8e+02; es 15; Conservative 0; Mismatches 3;
                          172 ACTACGAGTCCAAGGCAC 189
                                  18 ACTAGCATTCCAAGGCAC 1
                                                                        BP
                                                                       ADC78704 standard; DNA; 20
                                                                                                 (first entry)
                                                                                                 01-JAN-2004
                                                                                     ADC78704;
  Query Match
         Best Loca
Matches
                                                           RESULT 414
ADC78704
                                                                              à
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Diagnose of endometriosis in female involves detecting the presence of purified and isolated endometriotic haptoglobin and its functional analogs from patient sample. WPI; 2003-802186/75.

94US-00328451. 98US-00044604.

25-OCT-1994; 19-MAR-1998;

(TIMM/) TIMMS K L.

Timms KL;

27-NOV-2002; 2002US-00306903

US2003166014-A1.

Rattus sp.

04-SEP-2003

Example 7; SEQ ID NO 7; 28pp; English.

The invention describes a method of diagnosing endometriosis in a female suspected of having endometriosis comprising detecting the presence of a purified and isolated endometriotic haptoglobin (ENDO-1) and its functional analogues from a patient sample. The presence of the endometriotic haptoglobin is indicative of endometriosis. The invention provides purified and isolated glycoprotein and biologically functional analogues having specific physical and functional characteristics haptoglobin ENDO-1 cDNA.

Sequence 20 BP; 4 A; 6 C; 4 G; 6 T; 0 U; 0 Other;

Gape ö Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels

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ADD84533 standard; DNA; 20 ADD84533; RESULT 415
ADD84533/c
ID ADD845
XX
AC ADD845

Wed Apr 21 12:58:21 2004

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121P1F1 gene nested primer (NP) 2 SEQ ID NO:721.
 29-JAN-2004 (first entry)
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The present invention describes a composition (I) comprising a substance that modulates the status of 121PIF1 (gene and encoded protein), or a molecule that is modulated by 121PIF1, where the status of a cell that expresses 121PIF1 is modulated. The human 121PIF1 gene maps to chromosome 4q. (I) has cytostatic activity, and can be used in gene therapy, and in vaccines. The composition (I) can be used for diagnosing, preventing, prognosticating or treating patients with cancer that expresses 121PIF1, such as breast, colon, ovarian or lung cancer. The 12IPIF1 gene or its cragment can be used to elicit a humoral or cellular immune response. In 21PIF1 antibodies can be used in active or passive immunisation. 121PIF1 polymuclectides are useful as probes and primers for the amplification or detection of 121PIF1 polypeptides, or as tools for modulating or inhibiting the expression of 121PIF1 genes. The present sequence is used in the exemplification of 121PIF1 genes. 121P1F1; 121P1F1 modulation; human; chromosome 4q; cytostatic; gene therapy; vaccine; cancer; immune response; immunisation; primer; ss. Composition comprising a substance that modulates the status of 121PIF1, useful in diagnosing, preventing, prognosticating or treating patients with cancer that expresses 121PIF1, such as breast, colon, ovarian or 9 Challita-Eid PM, Hubert RS, Raitano AB, Faris M, Afar DEH, Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other; Example 1; Page 71; 285pp; English. 28-FEB-2002; 2002WO-US006242. 05-MAR-2001; 2001US-00799250. (AGEN-) AGENSYS INC. WPI; 2003-156757/15. WO200295009-A2. Homo sapiens. Jakobovits A; lung cancer, 28-NOV-2002

ö Gaps ch 3.1%; Score 13.2; DB 1; Length 20; l Similarity 83.3%; Pred. No. 3.8e+02; 15; Conservative 0; Mismatches 3; Indel8 Best Local Similarity Matches 15; Conserv

373 TCCTGGACCGCGACG 390 50 ò g

ADE65924 standard; DNA; 20 ADE65924; 

BP.

Human 161P2F10B protein-related PCR primer SegID36. (first entry)

(61P2F10B; cancer; cytostatic; gene therapy; vaccine; PCR; primer; ss;

A composition for diagnosing, preventing and treating cancer (e.g. prostatic, renal or uterine cancer) comprises 161P2F10B polynucleotides and polypeptides. Morrison KJM; Ge ₩ Jakobovits A, Raitano AB, Faris M, Hubert RS, Morrison RK, Challita-Eid PM; 07-NOV-2002; 2002WO-US036002. 07-NOV-2001; 2001US-00005480. 31-JAN-2002; 2002US-00062109. WPI; 2003-441560/41. (AGEN-) AGENSYS INC WO2003040340-A2 Homo sapiens. 15-MAY-2003 

This invention relates to a novel composition which comprises a substance that modulates the status of a novel protein (1612/2108) and its variants braving a sequence of 875 amino acids provided in the specification. The protein of the invention is over-expressed in certain cancers. The compounds of the invention may have cytostatic activity and the sequence of the Ispzings protein, and the gene which encodes it, may be useful for gene therapy or the development of a vaccine. The composition and methods of the invention are useful in diagnosing, preventing and treating ancount of a region of the gene encoding the human lifetation of a region of the gene encoding the human liPpzF10B protein during the exemplification of the resmall fileation of the invention.

Example 1; SEQ ID NO 36; 135pp; English

ô 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; ative 0; Mismatches 3; Indels Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other; Query Match Best Local Similarity 83.33 Matches 15, Conservative

373 TCCTGGACCGCGACG 390

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Gaps

ADD96944 Btandard; DNA; 20 (first entry) 29-JAN-2004 ADD96944; 417 RESULT 417 ADD96944/c 

BP.

Human protein 193P1E1B-related PCR primer SeqID59.

193PIE1B; tissue specific expression; cancer; cytostatic; gene therapy; cancer; human; PCR; RT-PCR; reverse transcription PCR; primer; ss.

Homo sapiens

WO2003050255-A2 19-JUN-2003. 06-DEC-2002; 2002WO-US039274.

07-DEC-2001; 2001US-00013312.

(AGEN-) AGENSYS INC

Hubert RS, Faris M, Challita-Eid PM, Raitano AB,

Jakobovits A;

rifampicin (RFP), streptomycin (SM), kanamycin (KM), isoniazid (INH) and ethambutol (EB). The rpoB gene is responsible for resistance to RFP; the responsible for resistance to SM and KM; the rpsL gene is responsible for resistance to SM, the inhA gene is responsible for resistance to INH; the katG gene is responsible for resistance to INH; and the embB gene is responsible for resistance to INH; invention also relates to nucleic acid probes having part of a nucleotide sequence of tubercle bacilli (TB) responsible for drug resistance and primers used to generate the probes. The present sequence is an oligonucleotide of the present invention. The oligonucleotides of the present invention can be used to enable the differentiation of drug resistance and the determination of infection with tubercle bacilli simultaneously

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WPI; 2003-532905/50.
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This invention relates to novel composition comprising a substance that modulates the status of a 433 residue protein, given in the specification with the DNA sequence encoding it, or a molecule that is modulated by the protein. The novel protein 193P121B exhibits tissue specific expression in normal adult tissue and is aberrantly expressed in certain cancers. Compositions which modulate the 193P121B protein may have cytostatic activity and the DNA sequence which encodes protein 193P121B may be useful in gene therapy. The composition of the invention may be useful primer which was used for the amplification of human 193P121B gene DNA during the exemplification of the invention.
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New composition comprising 193P1E18-related protein, useful for preventing or treating cancer.
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Pred. No. 3.8e+02;
); Mismatches 3; Indels
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                                                                                                    Example 1; SEQ ID NO 59; 260pp; English.
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0; Indels

3.1%; Score 13; DB 1; Loudo.0%; Pred. No. 2.6e+02; iive 0; Mismatches 0;

293 GGIGAAGGACCTG 305

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Local Similarity 100. les 13, Conservative

Best Loca Matches

Query Match

1 GGTGAAGGACCTG 13

Length 16;

Sequence 16 BP; 3 A; 4 C; 6 G; 3 T; 0 U; 0 Other;

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Wild-type capture oligonucleotide #13
373 TCCTGGACCGCGACGACG 390
                          m
                                                                                             AAF95086 standard; DNA; 16 BP
                          rccrcecceceaccace
                                                                                                                                                  (first entry)
                                                                                                                                                  23-MAY-2001
                                                                                                                       AAF95086;
                          20
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Ribozyme; erythropoietin; granulocyte colony stimulating factor;

interferon alpha; ss

WO200061729-A2. Homo sapiens.

19-OCT-2000.

Hammerhead ribozyme substrate #323.

16-FEB-2001 (first entry)

AAF02028;

AAF02028 standard; DNA; 17

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Tubercle bacillus, drug sensitivity, drug resistance, rifampicin, streptomycin, kanamycin, isoniazid, ethambutol, rpoB gene, rrs gene, rpsL gene, inhA gene, katG gene, embB gene, probe, PCR primer, ss.
                                                                                                                                                                                          Takenishi S;
                                                                                                                                                   99JP-00220357.
                                                                                                                                  02-AUG-2000; 2000EP-00306563
                                                                                    Mycobacterium tuberculosis
                                                                                                                                                                  (NISM ) NISSHINBO IND INC. (SYST-) SYSTEM RES INC.
                                                                                                                                                                                          Suzuki Y, Nishida M,
                                                                                                                                                                                                       WPI; 2001-246696/26.
                                                                                                                                                   03-AUG-1999;
                                                                                                    EP1076099-A2
                                                                                                                   14-FEB-2001.
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The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, BAR3/COUP-TF-1, the GATA transcription factor gene, IRP-2 and/or the CAATT Displacement Protein (CDF). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of Sequence 17 BP; 1 A; 9 C; 6 G; 1 T; 0 U; 0 Other; interferon alpha The present invention relates to oligonucleotides based on nucleotide sequences obtained from both wild-type tubercle bacilli (wrTB) that are susceptible to a drug and mutant-type tubercle bacilli (mrTB) that are resistant to a drug. The drugs used in the present invention are oligonucleotides, nucleic acid probes and primers are useful for ferentiating drug-resistance and determining infection with tubercle

Claim 21; Page 40; 114pp; English.

differentiating

sor genes, factor protein,

Enzymatic and antisense nucleic acid inhibition of repressor useful for producing e.g. granulocyte colony stimulating fact interferon alpha and erythropoietin.

Claim 37; Page 63; 164pp; English.

Mcswiggen J;

Blatt L, Zwick M, Pavco P,

WPI; 2000-647423/62.

11-APR-2000; 2000WO-US009721.

12-APR-1999; 99US-0129390P. (RIBO-) RIBOZYME PHARM INC. Length 17; 3.1%; Score 13; DB 1; Lt 100.0%; Pred. No. 2.9e+02; Query Match Best Local Similarity ö

Gaps

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BP.

Length 17;

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Human; POSH1 1; SH3 domain; POSH-like signalling protein 1; oncogene; Rho GTPase; signal transduction; gene expression; cancer; vaccine; gene therapy; transgenic; ss.
            Query Match 3.1%; Score 13; DB 1; Length 17; Best Local Similarity 100.0%; Pred. No. 2.9e+02; Matches 13; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                           Human POSHL1 scanning oligonucleotide SEQ ID NO 1752.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-JAN-2001; 2001MO-US000664.
30-JAN-2001; 2001MO-US000664.
30-JAN-2001; 2001MO-US000665.
30-JAN-2001; 2001MO-US000666.
30-JAN-2001; 2001MO-US000669.
30-JAN-2001; 2001MO-US000669.
30-JAN-2001; 2001MO-US000669.
30-JAN-2001; 2001US-US000670.
23-MAY-2001; 2001US-00864761.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-JAN-2002; 2002EP-00001165
                                                                                                                                                                                                                               ABV91039 standard; DNA; 17
                                                                                           338 CCAGGGCCGGCTG 350
                                                                                                                                                                                                                                                                                                          23-DEC-2002 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Shannon M;
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                                                                                                                                                                                            RESULT 42
ABV91039/
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                                                                                                                                                                                                                                                                                                                           Human; POSHL 1; SH3 domain; POSH-like signalling protein 1; oncogene; Rho GTPase; signal transduction; gene expression; cancer; vaccine; gene therapy; transgenic; ss.
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  Indels
                                                                                                                                                                                                                                                                                       Human POSHL1 scanning oligonucleotide SEQ ID NO 1749.
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  0; Mismatches
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30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.
23-MAX-2001; 2001WO-US0006670.
23-MAX-2001; 2001WS-00328205F.
                                                                                                                                                                         ABV91036 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28-JAN-2002; 2002EP-00001165.
                                     248 CCCGGGCTCGGCC 260
                                                                                                                                                                                                                                                    23-DEC-2002 (first entry)
  Matches 13; Conservative
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                                                                                                                                  RESULT 42, ABV91036, ABV91
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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (1), comprising a sequence of 730 amino acids (SI, ABB83999), a sequence having 65$ sequence identity to (SI), acids (SI, ABB83999), a sequence having 65$ sequence of 730 amino acids (SI) having 95$ devitations. especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene, foncedent for protein as as an adaptor protein that interacts with Rho family small Grasses as well as downstream components of the signal transduction pathway. (I) is useful conciding (II) for identifying a specific binding partner. (I) and nucleic acids (II) coused by altered expression of human POSHL1 including diagnosing and caused by altered expression of human POSHL1 including diagnosing and transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples of the invention. Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to
                                                                                          Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1.
                                                                                                                                                                                                                                                                                                                                Example 2; SEQ ID NO 1752; 60pp + Sequence Listing; English.
WPI; 2002-684061/74.
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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (1), comprising a sequence of 730 amino acids (S1, ABB83999), a sequence having 65% sequence identity to (S1), (S1) having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (1) and nucleic acids (II) encoding (1) are useful for diagnosing, monitoring disease and treating caused by altered expression of human POSHL1 including diagnosing and treating cancer, they useful in the development of vaccines and (II) is useful in gene therapy. (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonuclectide useful in examples of the invention. Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to Derwent by the European Patent Office

3equence 17 BP; 2 A; 7 C; 7 G; 1 T; 0 U; 0 Other;

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                                                                                                                                                                                                                                                                                  Human; POSHL 1; SH3 domain; POSH-like signalling protein 1; oncogene; Rho GTPase; signal transduction; gene expression; cancer; vaccine;
                                                                         Gaps
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                                            DB 1; Length 17; 2.9e+02;
                                                                     0; Indels
                                                                                                                                                                                                                                                             Human POSHL1 scanning oligonucleotide SEQ ID NO 1750.
                         Sequence 17 BP; 2 A; 7 C; 6 G; 2 T; 0 U; 0 Other;
                                               Query Match
3.1%; Score 13; DB
Best Local Similarity 100.0%; Pred. No. 2.9
Matches 13; Conservative 0; Mismatches
Derwent by the European Patent Office
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30-JAN-2001, 2001WO-US000664.
30-JAN-2001, 2001WO-US000666.
30-JAN-2001, 2001WO-US000666.
30-JAN-2001, 2001WO-US000667.
30-JAN-2001, 2001WO-US000669.
30-JAN-2001, 2001WO-US000669.
30-JAN-2001, 2001WO-US000669.
                                                                                                                                                                                   ABV91037 standard; DNA; 17 BP.
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                                                                                                338 CCAGGGCCGGCTG 350
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                                                                                                                                                           RESULT 422
ABV91037/c
ID ABV9103
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present sequence is that of a scanning oligonucleotide useful in examples of the invention. Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to Derwent by the Buropean Patent Office
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino acids (SI, ABBA399), a sequence having 6$$ sequence identity to (SI), (SI) having 95$ deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (I) and mucleic acids (II) encoding (I) are useful for diagnosing, monitoring disease and treating caused by altered expression of human POSHL1 including diagnosing and treating cancer, they useful in the development of vaccines and (II) is
                                                                                                                                                                                                                                                                                                         Gaps
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Rho GTPase; signal transduction, gene expression, cancer; vaccine,
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                                                                                                                                                                                                                                      Length 17;
                                                                                                                                                                                                                                                                                                  0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human POSHL1 scanning oligonucleotide SEQ ID NO 1751.
                                                                                                                                                                                                                                   Score 13; DB 1; Le
Pred. No. 2.9e+02;
0; Mismatches 0;
                                                                                                                                                                       Sequence 17 BP; 3 A; 7 C; 6 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                          3.1%; Scor.
100.0%; Pred
0; M
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30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US0006667.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.
23-MAY-2001; 2001WO-0S006770.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABV91038 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene therapy; transgenic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-JAN-2002; 2002EP-00001165
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABV91038;
                                                                                                                                                                                                                                          Query Match
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useful in gene therapy. (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples of the invention. Note: The present sequence did not form part of the printed specification, but is based on sequence difformation supplied to Derwent by the Buropean Patent Office

88888888

Sequence 17 BP; 3 A; 7 C; 6 G; 1 T; 0 U; 0 Other;

0; Gaps Query Match 3.1%; Score 13; DB 1; Length 17; Best Local Similarity 100.0%; Pred. No. 2.9e+02; Matches 13; Conservative 0; Mismatches 0; Indels 338 CCAGGGCCGGCTG 350 14 CCAGGGCCGGCTG 2 ઠે

ACC65163 standard; DNA; 17 BP. ACC65163; RESULT 424 ACC65163 

Murine oligonucleotide associated with tumour supression, SEQ ID 2410. 01-JUL-2003 (first entry)

Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; murine; tumour suppression; tumour reversion; apoptosis; virus resistance; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; ss.

Mus musculus.

WO2003025176-A2.

27-MAR-2003.

17-SEP-2002; 2002WO-IB004210.

17-SEP-2001; 2001FR-00011979.

(MOLE-) MOLECULAR ENGINES LAB

Telerman A, Amson R, Tuijnder M;

WPI; 2003-333167/31.

New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.

Disclosure; Page 312; 738pp; French.

The present invention relates to murine oligonucleotides (ACC62754-ACC68066), which are associated with tumour suppression, tumour reversion, apoptosis and virus resistance. The oligonucleotides are useful as (1) as probes and primers for detecting, identifying, quantifying and/or amplifying nucleic acid, e.g. as one component of a gene chip, in vitro as (anti) sense reagents; and (2) for production of recombinant polypeptides. The oligonucleotides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia

Sequence 17 BP; 3 A; 5 C; 7 G; 2 T; 0 U; 0 Other;

Gaps . 0 3.1%; Score 13; DB 1; Length 17; 100.0%; Pred. No. 2.9e+02; ive 0; Mismatches 0; Indels Best Local Similarity 100. Matches 13; Conservative

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AAA38383 standard; DNA; 18

AAA38383;

21-AUG-2000 (first entry)

Human Ets-2 phosphorothioate antisense oligonucleotide, SEQ ID NO:42.

Ets-2; human; transcription factor; chromosome 21q22.3; cancer; invasion; metastasis; skeletal abnormality; Down's syndrome; expression inhibition; phosphorothioate; antisense; ss.

Homo sapiens

US6054316-A.

25-APR-2000.

99US-00344579. 25-JUN-1999; 99US-00344579. 25-JUN-1999;

(ISIS-) ISIS PHARM INC.

Baker BF, Cowsert LM;

WPI; 2000-338495/29.

Antisense compound, 8-30 nucleobases in length, inhibiting the expression Ets-2 is useful for treating cancer and detecting Ets-2 expression.

Claim 3; Col 40; 31pp; English.

Sequences AAAAB349-A38388 represent antisense oligonuclectides targetted to the human Ets-2 gene, which inhibit its expression. The antisense coligonuclectides were designed to target different regions of the human Ets-2 RNA, and were analysed for their effect on Ets-2 mRNA levels by Ets-2 RNA, and were analysed for their effect on Ets-2 mRNA levels by C guantitative real-time PCR. The Ets-domain transcription factors are a family of proteins which are involved in controlling key cellular events condity of proteins hared by all members of this family. Through this condity members bind to the promoter regions of various genes at a GCA consensus sequence, thereby acting as either represents or at a GCA consensus sequence, thereby acting as either represents or activators of the gene. All but one Ets family protein bind to DNA as a conomer. Ets-2 has been implicated in the regulation of cellular conomer. Ets-2 has been implicated in the regulation of cellular chromosome 21q22.3 which is within a region known to undergo translocations associated with malignancies. Ets-2 has been found to be upregulated in several cancers, including lymphoblastic leukaemia. It may also play a role in the cancer phenotype, as it activates the urokinase cupsaminogen activator (UPA) promoter and the promoters of calmidation. High levels of turb and metalloproteinases are associated with tumour invasion and metastasis in breast cancers. As the Ets-2 gene is located on chromosome 21, which is triplicated in Down's syndrome, it is also thought to be responsible for the skeletal abnormalities present con this condition. The antisense oligonuclectides of the invention are useful for the treatment or prophylaxis of conditions associated with Ets-2 expression, especially cancer 

Sequence 18 BP; 3 A; 6 C; 4 G; 5 T; 0 U; 0 Other;

.. 0 Length 18; 0; Indels Query Match
3.1%; Score 13; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 13; Conservative 0; Mismatches 0;

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AAV46388-V46391 are PCR primers used in the isolation of a perchloroethane dehalogenase (PCE-DH) isolated from Dehalospirillum multivorans. This protein is used in a process for microbiological purification of water contaminated with chlorinated ethylenes and/or chlorinated propylenes. The process involves adding an electron donor a passing the water through a bloreactor containing a syntrophic mixed culture immobilised on a support, where the culture comprises at least one dehalogenating bacterium and at least one hydrogen-producing,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Microbiological purification of water contaminated with chlorinated olefin(s) - using combination of dehalogenating and hydrogen-producing
                                                                     Perchloroethane dehalogenase; PCE-DH; microbiological purification; water contamination; chlorinated ethylene; propylene; electron donor; bloreactor; dehalogenating bacterium; anaerobic microorganism;
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3.1%; Score 13; DB 1; Length 20;
Best Local Similarity 72.2%; Pred. No. 4.1e+02;
Matches 13; Conservative 1; Mismatches 4; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 4 A; 1 C; 8 G; 3 T; 0 U; 4 Other;
                                      D. multivorans PCE-Dehalogenase PCR primer #3.
                                                                                                                                                                                                                                 Location/Qualifiers
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/mcd_base= i
/note= "inosine"
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/mod_base= i
/note= "inosine"
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/note= "inosine"
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Sulfurospirillum multivorans.
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Eisenbeis M;
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18-NOV-1998
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                                                                                                                                  PCR primer;
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AAX21359/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to a method for detecting non-responders to anti-tumour necrosis factor (TNF) therapy. The method involves testing an individual for homozygosity for at least one single nucleotide polymorphism (SNP) in the gene coding for TNF receptor II, which is located on chromosome 1p36. Two novel SNPs, one in exon 2 (position 168 A/G) and one in exon 6 (position 587 T/G) which result in Lyssfelys and Met196Arg respectively, are also described. The method of the invention is useful for detecting non-responders to anti-TNF therapy such as infliximab therapy, or therapy of Crohn's disease. The genes containing the 2 novel polymorphisms are useful for diagnostic purposes in inflammatory, malignant or other chronic diseases. ABK3447-ABK33440 represent PCR primers used to amplify different regions of the human TNF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Detecting non-responders to anti-human necrosis factor therapy, comprises testing an individual for homozygosity for a single nucleotide polymorphism in the gene coding for the tumor necrosis factor receptor
                                                                                                                                                                                                                                                                                                 Human; anti-tumour necrosis factor receptor II; TNF receptor II; chromosome 1p36; infliximab therapy; Crohn's disease; malignant disorder; inflammatory disorder; chronic disease; receptor; PCR; primer; ss.
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                                                                                                                                                                                                                                                             Human TNF receptor II gene exon 4 PCR primer #2
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                                                                                                                                           ABK33430 standard; DNA; 19
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      CACTITCCIGGAC 380
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                           CACTTTCCTGGAC 16
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16-JAN-2002

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RESULT 426

ABK33430

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AAV46390;

AAV46390 ID AAV4 XX AC AAV4 XX

RESULT 427

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Query Match

Matches

Synthetic

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This sequence represents a preferred antisense oligomucleotide targetted against the gene encoding human microtubule-associated protein 4 (MAP4).

Inhibition of MAP4 expression was measured by determination of MAP4 mRNA levels in a variety of cell lines via real-time quantitative PCR. The cell lines used included the bladder carcinoma cell line PCR. The numan lung carcinoma cell line A549, human neonatal dermal fibroblasts and human embryonic keratinocytes. Microtubule-associated proteins comprise a group of proteins that mediate microtubule assembly and function which is required for cytoskeletal integrity. MAP4 is a member of the non-neuronal strabilising the microtubule lattice. MAP4 expression has been shown to be elevated in cells with mutant p53 oncogene expression, and is therefore linked to cancer chemcherapeutic drug sensitivity. These antisense molecules are useful for treating animals, particularly humans, having or being prone to a disease or condition associated with the expression of MAP4. The oligomucleotides are also useful for research and diagnostic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antisense oligonucleotides for inhibiting microtubule-associated protein 4 expression, useful in treating disorders associated with microtubule protein expression.
                                                                                                                                  methoxyethyl (2'-MOE) nucleotides"
methoxyethyl (2'-MOE) nucleotides"
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Best Local Similarity 100.0%; Pred. No. 4.1e+02;
Matches 13; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleotide sequence of PCR primer HCG-R2.
                                                                                                                   OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          primers AAX21358-X21359 were used to PCR amplify a fragment of the 17DE1 locus sequence as a control sequence for analysis of BAI gene expression in blots. The BAI genes (see AAX2135-X2137) are expressed specifically in the brain and play an important role in cancer formation in the brain. The BAI proteins can be used in drug compositions to diagnose, prevent or
                                                                                                                                                                                                                                                                                                                                                                                                                                                   New human BAI gene - is expressed in brain plays important role in cancer formation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Microtubule associated protein 4; MAP4; real-time quantitative PCR; expression; microtubule; assembly; function; cytoskeleton; structural; dynamic; stabilisation; lattice; overexpression; p53; oncogene; cancer; chemotherapy; tumour; drug sensitivity; antisense; therapy; hybridisation; inhibition; research; diagnostic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                               Human; BAII; brain; cancer; drug; diagnosis; prevention; treatment;
primer; PCR; amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human microtubule-associated protein 4 (MAP4) antisense oligo #37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Match 3.1%; Score 13; DB 1; Length 20; Local Similarity 100.0%; Pred. No. 4.1e+02; tes 13; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       'note = "Phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 2 A; 8 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 3; Page 16; 62pp; Japanese.
                                                                 Prime ElA for 17DE1 locus sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1. .5
/*tag= b
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ38502 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                  97JP-00176485.
                                                                                                                                                                                                                                                                                                                                     97JP-00150460.
                                                                                                                                                                                                                                                                                                                                                                        (SAKA ) OTSUKA PHARM CO LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  295 TGAAGGACCTGAG 307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                            21-MAY-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TGAAGGACCTGAG 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ..20
*tag=
                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1999-183823/16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  treat such cancers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Key
modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-FEB-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                         JP11032766-A.
                                                                                                                                                                                                                                                                                                                                       23-MAY-1997;
                                                                                                                                                                                                                                                                                                  16-JUN-1997;
                                                                                                                                                                                                                                                          09-FEB-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAZ38502;
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429

AAZ38502 RESULT

Query Match

Matches

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Gaps ö

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AAZ99396 standard; DNA; 20
                                                                                   Matches 13; Conservative
                                                                                 Best Local Similarity
                                                                                                            03-JUL-2000
                                                                                                        AAZ99396;
                                                                               Query Match
                                                                                                  AAZ99396/
ID AAZ9
                                                                                                 RESULT
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The specification describes a pre-trans-splicing molecule (PTW) which contains one or more target binding domains, a 3' splice region comprising a branch point, a pyrimidine tract and a 3' splice acceptor site, a spacer region separating the mRNA splice region from the target binding domain, and a nucleotide sequence to be trans-spliced. The method is used for the in vivo production of a trans-spliced molecule in a subset of cells. The PTM is used for producing chimeric mRNA molecule by contacting it with target pre mRNA which is useful for gene regulation, gene repair and targeted cell death particularly repair of cystic fibrosis trans-membrane regulator gene. The present primer was used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel pre-trans-splicing molecules for use in gene regulation, gene repair and targeted cell death particularly gene repair of cystic fibrosis trans-membrane regulator gene.
Pre-mRNA molecule; gene repair; pre-trans-splicing molecule; gene regulation; targeted cell death; cystic fibrosis trans-membrane regulator gene; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3.1%; Score 13; DB 1; Length 20; 100.0%; Pred. No. 4.1e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 3 A; 6 C; 9 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 6; Page 32; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Garcia-Blanco MA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (INTR-) INTRONN HOLDINGS LLC.
                                                                                                                                                                                                                                                                                                                                                                                       99WO-US018371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      98US-00133717
98US-00158863
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           the course of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2000-224360/19.
                                                                                                                                                                                                                                   WO200009734-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mitchell LG,
                                                                                                                                                        Unidentified
                                                                                                                                                                                                                                                                                                                                                                                           12-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13-AUG-1998;
23-SEP-1998;
                                                                                                                                                                                                                                                                                                         24-FEB-2000
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The specification describes a pre-trans-splicing molecule (PTM) which contains one or more target binding domains, a 3' splice region comprising a branch point, a pyrimidine tract and a 3' splice acceptor site, a spacer region separating the mRNA splice region from the target binding domain, and a nucleotide sequence to be trans-spliced. The method is used for the in vivo production of a trans-spliced molecule in a subset of cells. The PTM is used for producing chimeric mRNA molecule by contacting it with target pre mRNA which is useful for gene regulation, gene repair and targeted cell death particularly repair of cystic fibrosis trans-membrane regulator gene. The present primer was used to

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Query Match
3.1%; Score 13; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 4.1e+02;
Matches 13; Conservative 0; Mismatches 0; Indels

255 TCGGCCACGGTGC 267

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15

BP.

AAS00695 standard; DNA; 20

AAS00695/ RESULT

Gaps

;

255 TCGGCCACGGTGC 267

15 rcddcchcdcrdc 3

Sequence 20 BP; 3 A; 6 C; 9 G; 2 T; 0 U; 0 Other;

test a lacZ trans-splicing model

Novel pre-trans-splicing molecules for use in gene regulation, gene repair and targeted cell death particularly gene repair of cystic fibrosis trans-membrane regulator gene.

Garcia-Blanco MA;

Mitchell LG,

WPI; 2000-224360/19.

(INTR-) INTRONN HOLDINGS LLC.

98US-00133717. 98US-00158863.

13-AUG-1998; 23-SEP-1998;

Example 7; Page 42; 79pp; English.

##X#X#X#X#X###X

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Thymosin-beta-10-like protein; ephrin type-A receptor 8-like protein; 88; proteoglycan-like protein; fibromodulin; fibromectin; thymic immune cell; spermatogenesis; male infertility; neoplasia; red blood cell; platelet; small cell lung cancer; GPI-anchored ephrin-A ligand; prostate cancer; neurological disorder; cardiac disorder; vascular disorder; orthopaedic; inflammatory disease; rheumatoid archritis; connective tissue; congenital muscular dystrophy; chemotherapy; immunotherapy; PCR primer; BC 2.7.1.112.
                                                                              Forward PCR primer for analysis of ephrin type-A receptor 8-like protein.
                                                                                                                                                                                                                                                                                                                                                                     15-OCT-1999; 99US-0159805P.
18-OCT-1999; 99US-0159992P.
22-OCT-1999; 99US-0160952P.
12-OCT-2000; 2000US-00159805.
                                                                                                                                                                                                                                                                                                                                          13-OCT-2000; 2000WO-US028474
                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Taupier RJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CURA-) CURAGEN CORP
                                                                                                                                                                                                                                                                                 WO200129217-A2.
                                                      07-SEP-2001
                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Prayaga SK,
                                                                                                                                                                                                                                                                                                               26-APR-2001.
                       AAS00695;
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Pre-mRNA molecule; gene repair; pre-trans-splicing molecule; gene regulation; targeted cell death; lacZ; cystic fibrosis trans-membrane regulator gene; PCR primer; ss.

99WO-US018371

12-AUG-1999;

WO200009734-A2

Unidentified.

PCR primer HCG-R2 used to test a lacZ trans-splicing model

(first entry)

ВЪ.

New isolated polypeptides, NOV 1-3, having identity to thymosin-beta-10, ephrin type-A receptor 8 and proteoglycans, and polymucheotides, useful for treating male infertility, neurological or cardiac disease or rheumatoid arthritis.

Example 1; Page 83; 102pp; English.

The sequence represents a PCR primer used in expression analysis of ephrin type-A, receptor 8-like protein (NOV1). Thymosin-beta-10-like protein (NOV1), ephrin type-A receptor 8-like protein and proteoglycan[1] the proteins (NOV3) may be used in the diagnosis, treatment and proteoglycans. The protein of disorders caused by abnormal expression or activity of prevention of disorders caused by abnormal expression or activity of thymosin-beta-10, ephrin type-A receptor 8 and proteoglycans such as fibromedial and fibronectin. The polypeptides of the invention are useful in screening for agents that modulate their activity, and in determining predispositions to disorders. NOV1 is useful for treating conditions involving development, differentiation, and activation of thymic immune cells in pathologies related to spermatogenesis and male infertility, diagnosis of neoplasias, in diseases or pathologies of red blood cells or platelets, in detection of small cell lung cancer. NOV1 is useful for detecting cells expressing GPI-anchored chinched as a marker for prostate cancer, and in treating centuring and proteins are useful for treating orthopaedic disorders and/or injuries, and inflammatory diseases of connective tissues e.g. rheumatoid containing conservation or the prostate cancer, and intention contains are useful for treating orthopaedic disorders e.g. rheumatoid containing the conservation or the proposation of the contains are useful for treating orthopaedic disorders e.g. rheumatoid contains are useful for treating orthopaedic disorders e.g. rheumatoid contains are useful for treating orthopaedic disorders e.g. rheumatoid contains and inflammatory diseases of connective tissues e.g. rheumatoid contains are useful for treating orthopaedic disorders e.g. rheumatoid contains are useful for treating orthopaedic disorders e.g. rheumatoid contains are useful for treating orthopaedic disorders e.g. rheumatoid contains are useful for the contains are dysections of the contains are dysectined to the contains are dysecti

Sequence 20 BP; 4 A; 5 C; 6 G; 5 T; 0 U; 0 Other;

0; Gaps Match 3.1%; Score 13; DB 1; Length 20; Local Similarity 100.0%; Pred. No. 4.1e+02; tes 13; Conservative 0; Mismatches 0; Indels Query Match Best Loca Matches

7 GAGTGAAACTGCG 19

ઠે g

NOV2 cDNA specific forward PCR primer.

Membrane bound protein; secreted NOV protein; spermatogenesis; neoplasia; male infertility; angiogenesis; vascular pathology; orthopaedic disorder; inflammatory disease; congenital muscular distropy; muscular disorder; rheumatorid arthritis; fixed deformity; dysprothrombinaemia; disorder; arthrogryposis; hypoprothrombinaemia; hypokalaemic period paralysis; Smith-Lemil-Opitz syndrome; carcinoid tumour; centrocytic lymphoma; hypoprarathyroidism; Leigh syndrome; cervical carcinoma; leukaemia; macular dystrophy; vielliform type; McArdle disease; Meckel syndrome; multiple endocrine neoplasia I; multiple myeloma; hyperparathyroidism; parathyroid adenomatosis I; prolactinoma; digenic retinitis pigmentosa; somatotrophinoma; neovascular inflammatory vitreoretinopathy; arthritis; carcinoid syndrome; atopy; tendonitis; gene therapy; vaccine; PCR; RESULT 433
AAD38163/C
ID AAD38163 standard; DNA; 20 BP.
XX
AC AAD38163;
XX
DT 10-SEP-2002 (first entry)
XX
DT NOV2 CDNA specific forward PCR p;
XX
Membrane bound protein; secreted
XM
Membrane bound protein; fixed deform
XM
Machil-Lemli-Optiz syndrome; carc,
XM
Macular dystrophy; vitelliform I;
XM
Macarcinoid syndrome; neoplasia I;
XM
Mozocasopy; tendo
XX
XX
XX
XX
XX
NOCCASOSSOP79-A2.
XX
XX
XX
XX
XX
XX
10-OCT-2001; 2001WO-US031498.

12-OCT-2000; 2000US-00689486. 13-OCT-2000; 2000US-00687276. 09-OCT-2001; 2001US-00973424.

(CURA-) CURAGEN CORP

ä Bandaru Prayaga SK, Taupier RJ,

WPI; 2002-454545/48.

Novel membrane bound and secreted NOV polypeptides, for treating, diagnosing and preventing male infertility, neurological, cardiac and vascular pathologies, and inflammatory diseases e.g. rheumatoid arthritis.

Example 1; Page 118; 180pp; English.

The present invention relates to novel membrane bound and secreted NOV comported and polymucleotides encoding such proteins. Sequences of the invention are useful for treating or preventingly NOV-associated disorders of in humans and for manufacturing a medicament for treating a syndrome associated with human disease. They are useful for determining the presence of or predisposition to lung cancer. NOVI compounds are useful for development, differentiation and activation of thymic immune cells, pathologies related to spermatogenesis and male infertility, diagnosis of circulation such as red blood cells and platelets. NOVI nucleic acids are useful for detecting specific cell types and as specific marker for cancers in tissues. NOV2 and NOV4 compounds are useful for treating specific cell types and as specific marker for cancers in tissues. NOV2 and NOV4 compounds are useful to direct the cevelopment of nervous system and angiogenesis and for treating neurological, cardiac and vascular pathologies. NOV3 and NOV5 compounds concorned to treating various orthopsesis and NOV3 compounds concorned are useful for treating various orthopsesis of the marker for congenital muscular distormer, tissue e.g. rheumatoid arthritis, congenital muscular distormer, warious muscular discorders incorporate (for treating acopy, dysprothrombinaemia, hypoprothrombinaemia, type I and corporate (for treating acopy, dysprothrombinaemia, hypoprothrombinaemia, noveledent of lung, certical carcinoma, hyperparathyroidism, carcinoid syndrome, unltiple endocrine neoplasmial (multiple mysloms, parathyroid adenomatosis I, prolactinoma, hyperparathyroidism, carcinoid syndrome, and ultiple endocrine neoplasmial (multiple mysloms, parathyroid adenomatosis I, prolactinoma, hyperparathyroidism, carcinoid syndrome, and as vaccines (for the invention are also used in gene therapy and as vaccines I the exemplification of the invention

Sequence 20 BP; 4 A; 5 C; 6 G; 5 T; 0 U; 0 Other;

the invention

ö Query Match 3.1%; Score 13; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 4.1e+02; Matches 13; Conservative 0; Mismatches 0; Indels

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7 GAGTGAACTGCG 19

ò g

ABQ73441; ABQ73441/

(first entry) 02-OCT-2002

Human beta-chronic gonadotropin (HCG) RT-PCR primer HCG-R2.

Pre-trans-splicing molecule; PTM; spliceosome; cytostatic; gene therapy immunosuppressive; antimicrobial; gene regulation; gene repair; cancer; targeted cell death; genetic disorder; infectious disorder;

13 GAGTGAAACTGCG 1

ABQ73441 standard; DNA; 20 BP. RESULT 434

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Wed Apr 21 12:58:21 2004
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Novel cell having pre-trans-splicing molecules with target binding domains that target binding of PTM to pre-mRNA, 3' or 5' splice region, spacer region, nucleotide sequence to be trans-spliced to target-pre-mRNA.
                                                                                                                                     Garcia-Blanco MA, Baker CC, Puttaraju M;
Chao H;
autoimmune disease; proliferative disorder; PCR primer;
                                                                            08-JAN-2001; 2001US-00756095.

08-JAN-2001; 2001US-00756096.

08-JAN-2001; 2001US-00756097.

20-AFR-2001; 2001US-00841895.
                                                               08-JAN-2002; 2002WO-US000416
                                                                                                                        (INTR-) INTRONN INC.
                                                                                                                                                           WPI; 2002-566693/60.
                                   WO200253581-A2.
                                                                                                                                     Mitchell LG,
Mansfield GS,
             Homo sapiens.
Synthetic.
                                                 11-JUL-2002
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Example; Page 43; 229pp; English.

The present invention describes a cell (I) comprising pre-trans-splicing molecules (PTMs) (II) which have one or more target binding demains (IIa) controlled that target binding of PTM to pre-mRNA, 3 splice region (IIb) that includes branch point pyrimidine tract and 3'splice acceptor site, or 5' conprises (IIC), spacer region (IId) that separates RNA splice site (IIC), spacer region (IId) that separates RNA splice site (III) and (IIE); or (B) (IIC) (IId) and (IIE) be trans-comprised to target-pre-mRNA. Optionally, the cell comprises (II) either comprising: (A) (IIb) and (IIE); or (B) (IIC) (IId) and (IIE) The cell mmunosuppressive and antimicrobial activities, and can be used in gene therapy. (II) comprising one or more (preferably two or more) (IIB) and (III), (III) and (IIE), or (II) (III) and (IIE), or (III) (III) and (III), (III) comprises sequence encoding a toxin or translatable comprises sequence comprising components. The chimeric produced comprises sequences comprising exons 1-10 of cystic (II) that comprises nucleotide sequence regulator (CFIR). The chimeric RMA produced using (II) which either comprises (A) or (B) further comprises a nucleotide sequence regulator (CFIR). The chimeric RMA various diseases including genetic, infectious or autoimmune diseases and produced using dependence and to regulator comprises trans-membrane sonderer and to regulate gene expression in complication of the present invention

Sequence 20 BP; 3 A; 6 C; 9 G; 2 T; 0 U; 0 Other;

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Query Match
3.1%; Score 13; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 4.1e+02;
Matches 13; Conservative 0; Mismatches 0; Indels
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255 TCGGCCACGGTGC 267 |||||||||||||||||||1|||1|||15 TCGGCCACGGTGC 3

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RESULT 435 ABQ73457/c ID ABQ73457 standard; DNA; 20 X XX

(first entry) 02-OCT-2002 ABQ73457;

rng.res

Human beta-chronic gonadotropin (HCG) related PCR primer HCG-R2

Pre-trans-splicing molecule, PTM; spliceosome; cytostatic; gene immunosuppressive; antimidrobbal; gene regulation; gene repair; targeted call death; genetic disorder; infectious disorder; autoimmune disease; proliferative disorder; FCR primer; 89.

Homo sapiens Synthetic.

WO200253581-A2.

08-JAN-2002; 2002WO-US000416 11-JUL-2002.

08-JAN-2001; 2001US-00756095. 08-JAN-2001; 2001US-00756096. 08-JAN-2001; 2001US-00756097. 20-ARR-2001; 2001US-00841492.

(INTR-) INTRONN INC.

Garcia-Blanco MA, Baker CC, Puttaraju M; Chao H; Mitchell LG, Mansfield GS,

WPI; 2002-566693/60.

Novel cell having pre-trans-splicing molecules with target binding domains that target binding of PTM to pre-mRNA, 3' or 5' splice region, spacer region, nucleotide sequence to be trans-spliced to target-pre-

Example, Page 53; 229pp; English.

Sequence 20 BP; 3 A; 6 C; 9 G; 2 T; 0 U; 0 Other;

exemplification of the present invention

Query Match 3.1%; Score 13; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 4.1e+02; Matches 13; Conservative 0; Mismatches 0; Indels

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Wed Apr 21 12:58:21 2004
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255 TCGGCCACGGTGC 267 ||||||||||||||||||1|| 15 TCGGCCACGGTGC 3

ઠે g ABL43850

ABL43850 standard; DNA; 16 BP ABL43850;

11-APR-2002 (first entry)

Human chromosome 1p36-35 PCR primer SEQ ID NO:894.

chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome; PCR primer; ss. Human;

Homo sapiens.

JP2001321190-A.

20-NOV-2001.

12-MAR-2001; 2001JP-00068285.

10-MAR-2000; 2000JP-00066716

(RIKA ) RIKAGAKU KENKYUSHO. (GENO-) GENOTEX YG.

WPI; 2002-144136/19.

Arraying genome clones.

Claim 4; Page 22; 528pp; Japanese.

The present invention describes a method of arraying genome clones. The method comprises: (a) clones of the genomic libraries contained in method comprises: (a) clones of the genomic libraries contained in multiwell plates numbered for discrimination are mixed in each of the multiwell plates; (b) a primer designed based on the chromosome marker sequence is added to the mixture to carry out an amplification reaction; (c) a signal corresponding to the marker is detected from the resultant amplified product to specify the discrimination Nos. of the multiwell plates containing the clones having said marker sequence; (d) the order of the maximum in the specified discrimination Nos. to array the multiwell containination Nos. that the same discrimination Nos succeed to containination Nos. the miltiwell plates of the specified discrimination Nos. to array the multiwell containination Nos. are mixed respectively in each wells of longitudinal and lateral directions; (f) the mixed clones are cultured and the call and lateral directions; (f) the mixed clones are cultured and the reconstituted as the positions on the chromosome and arrayed. The clones are processed to plates are specified from the edsected result; and (i) the clones are reconstituted as the positions on the chromosome and arrayed. The mixed plates for human chromosome 1936-35 DNA, and ABL43323 to ABL43534 to represent PCR primers for human chromosome 21q22.1, which are specifically claimed for use in the present invention 

Sequence 16 BP; 3 A; 7 C; 3 G; 3 T; 0 U; 0 Other;

Gaps . 0 Query Match

3.0%; Score 12.8; DB 1; Length 16;
Best Local Similarity 87.5%; Pred. No. 2.8e+02;
Matches 14; Conservative 0; Mismatches 2; Indels

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44 TGGCCACCACTCAGAG 59

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1 TGGCCCCCACTCATAG 16

RESULT 437 AAQ47599/c ID AAQ47599 standard; cDNA to mRNA; 17 BP.

(revised)
(first entry) 25-MAR-2003 26-JAN-1994 AAQ47599;

7 . FITT

Rat C RATRJG9/B-1258 jun-B specific probe.

Probe; quantification; human; GTP binding protein; G protein; alpha subunit; specific mRNA; detection; hybridisation; diagnosis; pathophysiology; disease state; hereditary; cancer; infectious; osteodystrophy; pituitary tumour; acromegaly; melanoma cells; diabetes; PCR; polymerase chain reaction; ss.

Synthetic.

WO9315221-A1.

05-AUG-1993.

93WO-US000977 29-JAN-1993;

29-JAN-1992; 24-MAR-1992; 12-NOV-1992;

92US-00827208. 92US-00857059. 92US-00974409.

Akitaya T, Cooper A, Mitsuhashi M; (HITB ) HITACHI CHEM CO LTD. (HITB ) HITACHI CHEM RES CENT INC.

WPI; 1993-258695/32.

Quantitating messenger RNA in sample - using immobilised-polynucleotide having sequence complementary to sequence unique to the MRNA.

Example 9, Page 71; 177pp; English.

The sequences given in AAQ47594-603 show regions of homology between jun sequences and the jun-B specific probe B-1258 which may be of use as jun-B specific probes. They were used in the method of the invention for the gracific probes. They were used in the method of the invention for the guzify the many from cells. The claimed method comprises identifying a purify the mRNA from cells. The claimed method comprises identifying a polymucleotide sequence unique to the mRNA, and immobilising an oligomer componente are washed from the support such that the unique sequence then incubated with the insoluble support such that the unique sequence components are washed from the support and bound RNA is labelled in such components are washed from the support. The amount of bound label is then a way that the label is incorporated onto the support relative to the amount of mRNA on the support. The amount of bound label is fine cell of the amount of many be used for the reliable, rapid, simultaneous cell agnosing and recognition of pathophysiclogy of various disease states, cancer, and infectious disease states, cancer, and infectious disease states, cancer, and infectious disease states. Cell deficiency of Gs protein is the molecular basis of hereditary contain mutant ds proteins. G proteins are also involved in invasive contain mutant ds proteins. G proteins are also involved in invasive contain mutant ds proteins are also involved in invasive contain metastatic melanoma cells, and diabetes. See also AAQ473B1-666. 

Sequence 17 BP; 3 A; 8 C; 4 G; 2 T; 0 U; 0 Other;

Gaps ô Query Match
3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels

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142 TGGCGGTGGAGGCCGG 157 16 TGGCGGTGGACGCCAG

ઠે 셤 RESULT 438

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AAT50887-T50904 represent oligomucleotides of the invention. These sequences are all probes for interleukin-6 receptor (IL-6R) mRNA. IL-6 is one of the most well characterised of the cytokines. It functions through interacting with at least two transmembrane glycoprotein receptor molecules on the surface of target cells. The receptors are the IL-6R, and the signal transductor py IL-6 involves the concerted action of both IL-6R and gpl30. IL-6 overproduction is implicated in many different disease states, particularly in cellular proliferation associated with these diseases. These sequences bind to the IL-6R coding sequence, thereby inhibiting IL-6R production. The sequences therefore inhibit the functioning of IL-6R production. The coligomucleotides are especially useful for treating cancer (e.g. renal cell carcinoma), autoimmune diseases or viral infections. They can also be used as probes for detecting IL-6 receptor mRNA, especially for evaluating the effectiveness of drugs in reducing IL-6 receptor mRNA
                                                                                                                                               Probe; interleukin-6 receptor; IL-6R; cytokine; cellular proliferation; transmembrane glycoprotein receptor; signal transducer; gp130; inhibitor; IL-6; cancer; renal cell carcinoma; autoimmune disease; viral infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oligo:nucleotide(s) complementary to interleukin-6 receptor mRNA - for treating proliferative diseases, e.g. cancer, auto-immune diseases or viral infections.
                                                                                                                                                                                                                                                                                      1. .17
/*tag= a
/note= "optionally phosphorothioated"
                                                                                                                  Probe #3 for interleukin-6 receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Dattagupta N, Naidu YM;
                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 16; 18pp; English.
AAT50889/c
ID AAT50889 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         95US-00484666.
95US-00486408.
                                                                                                                                                                                                                                                                                                                                                                                                                         96EP-00304315.
                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GENP-) GEN-PROBE INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1997-023093/03.
                                                                                                                                                                                                                                                                                                                                                                                                                         07-JUN-1996;
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                                                                                                                                                                                                                                                                                      misc_feature
                                                                                                                                                                                                                                                                                                                                                                                         11-DEC-1996.
                                                                                  26-AUG-1997
                                                                                                                                                                                                                                                                                                                                                       EP747386-A2
                                                                                                                                                                                                       therapy; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Brown SJ,
                                                                                                                                                                                                                                     Synthetic
                                                  AAT50889;
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3.0%; Score 12.8; DB 1; Length 17;
87.5%; Pred. No. 3.2e+02;
ative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                               Oligo #13 used to isolate human chromosome 16 sequences.
                                                                                                                                                             86.
                                                            305 GAGCCCCGGGGACCGC 320
                                                                                                                                                             AAT85503 standard; cDNA; 17
                                                                                17 GAGCCCGGAGCCCGC 2
                                                                                                                                                                                                                   (first entry)
                                 Matches 14; Conservative
                   Similarity
                                                                                                                                                                                                                   17-NOV-1997
                                                                                                                                                                                          AAT85503;
    Query Match
Best Local &
                                                                                                                                  RESULT 440
                                                                                                                                               AAT85503
ID AAT8
XX AAT8
XX AAT8
XX IT-N
DT 17-N
DE Olig
XX Huma
KW AUGM
                                                              δ
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                                                                                                                                                              3.0%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 3.2e+02; tive 0; Mismatches 2; Indels
                                                                                                                                   Sequence 17 BP; 3 A; 7 C; 4 G; 3 T; 0 U; 0 Other;
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CGAGGGCCGCGCAGTG 89

74 17

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Conservative

14;

Best Loca Matches

Local Similarity

Query Match

ceaeceacreceaere 2

RESULT 439 AAX68712/c

Human; netrin; ATPase binding cassette transporter; ribosomal L3; augmenter of liver regeneration; hNET; hABC3; SEM L3; hALR; chromosome 16; exon trapping; axon; chicken; laminin domain; C. elegans;

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The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fms-like tyxosine kinase 1 (ELL1), kinase insert domain containing receptor (XDR) and/or foetal liver kinase 1 (Elk-1) (e.g. tumour anglogenesis, coular diseases, postriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAXX7275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention

Seguence 17 BP; 1 A; 6 C; 8 G; 0 T; 2 U; 0 Other;

Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.

Claim 4; Page 46; 218pp; English.

Stinchcomb D, Escobedo J;

95US-0005974P. 96WO-US017480,

25-OCT-1996; 26-OCT-1995; 11-JAN-1996;

01-MAY-1997.

(RIBO-) RIBOZYME PHARM INC. (CHIR ) CHIRON CORP.

Pavco P, Mcswiggen J, WPI; 1997-259017/23.

Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like, tyrosine kinase 1; kinase insert domain containing receptor;

foetal liver kinase 1; ss.

Homo sapiens. WO9715662-A2.

Human flt1 VEGF receptor hammerhead ribozyme substrate #7.

(first entry)

28-JUL-1999

AAX68712;

BP.

AAX68712 standard; RNA; 17

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Claim 34; Page 62; 98pp; English
(GENZ ) GENZYME CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         regeneration.
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Van Raay TJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23-JAN-1997.
                    Landes GM, I
                                                                                                                                                                                                                                                                                                                                                                                                                                          17-NOV-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                       AAT85480;
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                                                                                                                                                                                                                                                                                                                                                                                         AAT85480
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                                                                                                                                                                                                                                                           The sequences given in AAT85503-06 are oligos which were used in the isolation of coding sequences from human chromosome 16. The invention contains details of the sequences encoding human netrin (hNET), human ATP828 Binding Cassette transporter (hABG2), human ribosomal L3 (SEM L3), and human augmenter of liver regeneration (hARR). The hNET gene can be used to develop chemostractants for use in axon regeneration. The hABG3 gene may be used in therapeutic applications for cystic fibrosis. The hABC3 gene may be used to develop products for treating damaged liver and liver diseases. The products can also be used for detection, diagnosis and screening assays. These oligonucleotides of may be used as primers in exon trap amplification experiments
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; netrin; ATPase binding cassette transporter; ribosomal L3; augmenter of liver regeneration; hNET; hABC3; SEM L3; hALR; chromosome 16; exon trapping; axon; chicken; laminin domain; C. elegans; UNC-6; cystic fibrosis; ss.
                                                                                                                                                                                                 New isolated human chromosome 16 genes - encode netrin, ATPase binding cassette transporter, ribosomal L3 sub-type or augmenter of liver
                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Gaps
                                                                                                                                                 Connors TD, Dackowski WR, Klinger KW;
                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17 BP; 4 A; 3 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oligo #1 hybridises to hABC3 cDNA sequence.
                                                                                                                                                                                                                                            Disclosure, Page 18; 98pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAT85475 standard; cDNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                       288 AAGCTGGTGAAGGACC 303
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                                                                                   96WO-US010469.
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                                                                                                        95US-0000596P
  UNC-6; cystic fibrosis; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                 Burn IC,
                                                                                                                            (GENZ ) GENZYME CORP.
                                                                                                                                                                               WPI; 1997-108959/10.
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                                                                                                                                                                                                                         regeneration.
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                                                                                     17-JUN-1996;
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                                                                                                                                                 Landes GM, 1
Van Raay TJ;
                                          WO9702346-A2
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                                                               23-JAN-1997
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                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT85475;
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The sequences given in AAPBS475-83 hybridise under stringent conditions to the sequence encoding the ATPase binding cassette transporter protein (habs3). The hABG3 genomic sequence was isolated from human chromosome 16 by exon trapping, hABG3 cDNA contains an open reading frame of 1685 amino acids. Comparison of ABC1, ABC2 and hABG3 reveals significant conservation in the regions surrounding the two ATP binding cassettes. The ATP binding cassettes of hABG3 flank a large linker domain containing numerous polar residues. The presence of these features in the linker domain suggests that this domain may play a regulatory role similar to the R domain of CPTR. The hABC3 gene may be used in therapeutic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; netrin; ATPase binding cassette transporter; ribosomal L3; augmenter of liver regeneration; hNET; hABC3; SEM L3; hALR; chromosome 16; exon trapping; axon; chicken; laminin domain; C. elegans; UNC-6; cystic fibrosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New isolated human chromosome 16 genes - encode netrin, ATPase binding cassette transporter, ribosomal L3 sub-type or augmenter of liver
                                                                                                                                           New isolated human chromosome 16 genes - encode netrin, ATPase binding cassette transporter, ribosomal L3 sub-type or augmenter of liver regeneration.
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Dackowski WR, Klinger KW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17 BP; 4 A; 3 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oligo #6 hybridises to hABC3 cDNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Burn IC, Connors ID,
       Connors TD,
                                                                                                                                                                                                                                                                                                                    Claim 29; Page 61; 98pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAT85480 standard; cDNA; 17 BP.
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       Burn TC,
                                                                                                             WPI; 1997-108959/10.
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The sequences given in AAT85475-83 hybridise under stringent conditions to the sequence encoding the ATF8se binding casette transporter protein (hABC3). The hABC3 genomic sequence was isolated from human chromosome 16 by exon trapping. hABC3 cDNA contains an open reading frame of 1685 amino acids. Comparison of ABC1, ABC2 and hABC3 reveals significant conservation in the regions surrounding the two ATF binding casettes. The ATF binding casettes of hABC3 flank a large linker domain containing numerous polar residues. The presence of these features in the linker the ATF domain may play a regulatory role similar to the R domain of CFTR. The hABC3 gene may be used in therapeutic *6666666666668*

Query Match
3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels Sequence 17 BP; 4 A; 3 C; 8 G; 2 T; 0 U; 0 Other;

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RESULT 443 AAV95292/c ID AAV95292 standard; RNA; 17 BP.

AAV95292;

24-FEB-1999 (first entry)

Human c-fos target sequence nucleotide position 268.

Human; c-fos; hammerhead ribozyme; hairpin ribozyme; target site; cancer; oncogene; leukaemia; neuroblastoma; diagnosis; genetic drift; mutation; diseased cell; ss.

Homo sapiens

WO9832846-A2.

30-JUL-1998

20-JAN-1998;

97US-0037658P. 97US-00998099. 23-JAN-1997; 24-DEC-1997;

(RIBO-) RIBOZYME PHARM INC.

WPI; 1998-427942/36.

Jarvis T, Mcswiggen JA, Stinchcomb DT;

Enzymatic nucleic acid molecules which specifically cleave RNA derived from a c-fos gene - useful for treating conditions related to levels of c-fos, especially cancer.

Claim 2, Page 50, 72pp, English.

The present invention describes an enzymatic nucleic acid molecule which specifically cleaves RNA derived from a c-fos gene. AAV95401 to AAV95540 and AAV95541 to AAV95541 represent hammerhead ribozymes and hairpin ribozymes, respectively, which specifically cleave human c-fos. AAV95261 to AAV95400 and AAV95585 to AAV95628 represent human c-fos target cancer associated with elevated levels of c-fos oncogene, especially leukaemias, neuroblastomas and lung, breast and colon cancers. The ribozymes may also be used a diagnostic tools to examine genetic drift and mutations within diseased cells, or to detect the presence of c-fos RNA in a cell

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Gaps
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0
Query Match 3.0%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 3.2e+02; Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                        286 CCAAGCTGGTGAAGGA 301
                                                                                                                               17 ccardcredadadeda 2
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Sequence 17 BP; 2 A; 7 C; 3 G; 0 T; 5 U; 0 Other;

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RESULT 444 AAV45792

AAV45792 standard; DNA; 17 BP.

AAV45792;

24-NOV-1998 (first entry)

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Gaps ö Primer NONA PCR-R.

Gene bank; combinatorial library; phagemid display; phage display; cosmixplexing; receptor; ligand; autoimmune disease; ss.

Synthetic.

WO9833901-A2

06-AUG-1998.

98WO-EP000533. 02-FEB-1998;

97EP-00101539 31-JAN-1997; (COSM-) COSMIX MOLECULAR BIOLOGICALS GMBH.

Collins J, Roettgen P;

WPI; 1998-437456/37.

Banks containing genes with restriction enzyme sites that generate specific cohesive ends - allowing production of large phage or phagemid display libraries, for screening to identify ligands for medical, diagnostic etc. use.

Example 1; Page 45; 87pp; English.

In a cosmiplexing method of the invention for the generation of double-stranded DNA inserts, the single-stranded hypervariable DNA oligos NONA-CA, NONA-CA, MONA-CA, 

Sequence 17 BP; 3 A; 4 C; 7 G; 3 T; 0 U; 0 Other;

0 Match 3.0%; Score 12.8; DB 1; Length 17; Local Similarity 87.5%; Pred. No. 3.2e+02; les 14; Conservative 0; Mismatches 2; Indels Query Match Best Local S

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Gaps

262 CGGTGCACCTGGAGCA 277

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CGGGGTACCTGGAGCA 17

rng.res

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Oligonucleotides AAV16310-25 are used to clone addional sequences of nucleic acids encoding human netrin (hNET), human ATP binding cassette transporter (LABC2), human ribosomal 13 (RPL31), and human augmenter of liver regeneration (hALR). Partial DNA sequences from these genes were isolated using exon traps AAW46753-57. Genetrapper, 3' RACE and RT-PCR were employed to identify additional sequences. The antisense oligonucleotides of the isolated sequences are used to modulate expression of hNET, hABC3, RPL31 or hALR, and prevent its translation. Antibodies against hNET, hABC3, RPL31 and hALR can be used to block binding of their naturally occurring ligands. The host cells containing vectors with DNA inserts encoding the proteins can be used to a method for identifying compounds which bind to hNET, hABC3, RPL31 or hALR.

Modulation or alteration of hABC3 substrate specificity may have significant therapeutic implications for cystic fibrosis. hALR could be used in the treatment of damaged liver
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human chromosome 16 genes encoding netrin, ATP binding cassette transporter, ribosomal L3 and augmenter of liver regeneration proteins useful for, e.g. treatment of liver disease and cystic fibrosis.
                                                                                                                                                                         Human; netrin; hNET; ATP binding cassette transporter; hABC3; ribosomal L3; RPL3L; augmenter of liver regeneration; hALR; treatment; trapping; modulation; expression; antibody; identification; binding; substrate specificity; ligand; exon trap; PCR primer; amplify; ss.
                                                                                                                                      Primer used to clone additional sequences of ABC, NET, ALR and RPL3L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Connors TD, Dackowski WR, Van Raay TJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17 BP; 4 A; 3 C; 8 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                       96US-00665259.
96US-00720614.
96US-00762500.
                                   AAV16316 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                      97WO-US000785.
                                                                                                         03-JUN-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Burn TC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GENZ ) GENZYME CORP.
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01-OCT-1996;
09-DEC-1996;
                                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                WO9748797-A1.
                                                                                                                                                                                                                                                                                                                                                                                                      16-JAN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Landes GM,
Klinger KW;
                                                                                                                                                                                                                                                                                                                                                                   24-DEC-1997
                                                                                                                                                                                                                                                                       Synthetic
                                                                      AAV16316;
RESULT 445
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Gaps ö Query Match
3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels

288 AAGCTGGTGAAGGACC 303

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AAV16329 standard; DNA; 17 AAV16329; RESULT 446 AAV16329 ID AAV1 XX AC AAV1

03-JUN-1998 (first entry)

Primer used to clone additional sequences from human ABC3.

Human, ATP binding cassette transporter, hABC3; cystic fibrosis; treatment, trapping; modulation; expression; antibody; identification; binding; substrate specificity; ligand; exon trap; PCR primer; amplify;

Synthetic.

Homo sapiens.

WO9748797-A1

24-DEC-1997.

97WO-US000785. 16-JAN-1997;

96US-00665259. 96US-00720614. 96US-00762500. 17-JUN-1996; 01-OCT-1996; 09-DEC-1996;

(GENZ ) GENZYME CORP.

Dackowski WR, Van Raay TJ; Burn TC, Connors TD,

Landes GM, Klinger KW;

WPI; 1998-063138/06.

Human chromosome 16 genes encoding netrin, ATP binding cassette transporter, ribosomal L3 and augmenter of liver regeneration proteins useful for, e.g. treatment of liver disease and cystic fibrosis.

Claim 40; Page 25; 220pp; English.

oligonucleotides AAV16326-32 are used to clone addional sequences from nucleic acids encoding human ATP binding cassette transporter (hABC3).

Partial DNA sequences from the genes was isolated using exon traps additional bequences. The ABC gene is located in the PKD1 locus, between additional sequences. The ABC gene is located in the PKD1 locus, between the ECM1 and D162291 markers in a centromeric to telomeric orientation.

The sequence shows homology with murine ABC1 and ABC2 genes. The ABC proteins are responsible for the transport of a wide variety of substrates across cell membranes. Proteins in this family are linked by strong structural similarities. ABC transporters govern unidirectional cransport of molecules into or out of cells and across subcellular membranes. The antisense oligonucleotides of the ABC3 gene sequence are membranes. The antisense oligonucleotides of the ABC3 gene sequence are commented to be used to block binding of its naturally occurring against ABC can be used to block binding of its naturally occurring cortein can be used in a method for identifying compounds which bind to be significant therapeutic implications for cystic fibrosis 

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Sequence 17 BP; 4 A; 3 C; 8 G; 2 T; 0 U; 0 Other;

ö Query Match
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels

288 AAGCTGGTGAAGGACC 303 

RESULT 447

AAA364116
ID AAA36411 standard; DNA; 17 BE
XX
AC AAA36411;
AC AAA36411;
DT 26-JUL-2000 (first entry)

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A method has been developed for detecting the presence or absence of a single nucleotide polymorphism (SNP) allele in a genomic sample. The method comprises preparing a reduced complexity genome (RGG) from the genomic sample and analysing the RGG for the presence or absence of a SNP allele. The method can be used to characterise a tumour, to generate a genomic pattern for an individual genome or to generate a genomic classification code for a genome. The method can be used to assess whether a subject is at risk for developing a disease or to identify a set of SNP alleles associated with a disease. The method can also be used to perform linkage analysis. AAA35944 to AAA35947 represent sequences used in the exemplification of the present invention. AAA35948 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Detection of single nucleotide polymorphisms in genomes by preparation and analysis of reduced complexity genomes, useful for genotyping, fingerprinting and determining allele frequency of SNPs.
                                                   Human; single nucleotide polymorphism; SNP; genotyping; DNA analysis; allele specific ollogonucleotide; ASO; reduced complexity genome; RCG; genomic classification; identification; by fingerprinting; tumour characterisation; hybridisation; 88.
                Human genomic SNP allele specific oligonucleotide SEQ ID NO:477
                                                                                                                                                                                                                                                                                                                                                                                 Charest A;
                                                                                                                                                                                                                                                                                                                                            (MASI ) MASSACHUSETTS INST TECHNOLOGY
                                                                                                                                                                                                                                                                                                                                                                                 Housman DE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 67; 111pp; English.
                                                                                                                                                                                                                                                                   99WO-US022283
                                                                                                                                                                                                                                                                                                     98US-0101757P
                                                                                                                                                                                                                                                                                                                                                                                   Landers JE, Jordan B,
                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-293181/25.
                                                                                                                                                                                       WO200018960-A2.
                                                                                                                                                                                                                                                                                                       25-SEP-1998;
                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                   24-SEP-1999;
                                                                                                                                                                                                                              06-APR-2000
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ö Length 17; 2; Indels Sequence 17 BP; 1 A; 7 C; 2 G; 7 T; 0 U; 0 Other; 3.0%; Score 12.8; DB 1; 87.5%; Pred. No. 3.2e+02; tive 0; Mismatches 2; Local Similarity 87.5 es 14; Conservative Query Match Best Loca Matches

204 GTGAAAGCAGAGACT 219 17 GAGAAAGCAGAGGACT 2 d ઠે

AAF02688 standard; DNA; 17 BP. AAF02688; RESULT 448 AAF02688/c 

(first entry) 16-FEB-2001

Hammerhead ribozyme substrate #983.

Ribozyme; erythropoietin; granulocyte colony stimulating factor; interferon alpha; ss.

Homo sapiens

WO200061729-A2

19-0CT-2000

11-APR-2000; 2000WO-US009721

99US-0129390P 12-APR-1999;

(RIBO-) RIBOZYME PHARM INC.

Mcswiggen J; Blatt L, Zwick M, Pavco P,

Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin. WPI; 2000-647423/62. 

Claim 37; Page 78; 164pp; English.

The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor. EAR3/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the GAAT Displacement Protein (CPP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoietin, granulocyte colony stimulating factor protein and interferon alpha

Sequence 17 BP; 0 A; 9 C; 4 G; 4 T; 0 U; 0 Other;

Gaps . 0 Length 17; 2; Indels 7 Match 3.0%; Score 12.8; DB 1; Local Similarity 87.5%; Pred. No. 3.2e+02; nes 14; Conservative 0; Mismatches 2; Query Match Best Loca Matches

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AAF05332 standard; DNA; 17 BP. RESULT 449 

(first entry) 16-FEB-2001 AAF05332;

Hammerhead ribozyme substrate #2551.

Ribozyme; erythropoietin; granulocyte colony stimulating factor; interferon alpha; ss.

Homo sapiens

WO200061729-A2. 19-OCT-2000.

11-APR-2000; 2000WO-US009721.

99US-0129390P (RIBO-) RIBOZYME PHARM INC. 12-APR-1999;

Mcswiggen J; Blatt L, Zwick M, Pavco P,

WPI; 2000-647423/62.

Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoletin.

Claim 18; Page 114; 164pp; English

The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes

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88888888

(first entry)

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Universal array of oligonucleotides tags attached to a solid substrate along with locus-specific tagged oligonucleotides useful in genotyping using single base extension reactions.
                                                                                                                                                                                 Oligonucleotide array; genotyping; single base extension reaction; PCR primer; polymorphic locus; single nucleotide polymorphism; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hirschhorn JN, Huang X, Kaplan P, Lander ES,
                                                                                                                                              Reverse primer #67 used in multiplexing PCR/SBE assay
                                                                                                                                                                                                                                                                                                                                                                                                                                                         (WHED ) WHITEHEAD INST BIOMEDICAL RES. (AFFY-) AFFYMETRIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 7; Page 55; 70pp; English.
                                    AAC73338 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                           27-MAR-2000; 2000WO-US008069.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-656171/63.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fan J, Hirschhorn
Ryder T, Sklar P;
                                                                                                                                                                                         Oligonucleotide
                                                                                                                                                                                                                                                                                    WO200058516-A2.
                                                                                                                                                                                                                                                                                                                                                                                                  26-MAR-1999;
23-JUN-1999;
                                                                                                                                                                                                                                                Unidentified.
                                                                                                               02-FEB-2001
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                                                                           AAC73338;
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  RESULT 451
AAC73338
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 orphan receptor, EAR3/COUP-TF-1, the GATA transcription factor gene, IRR-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoietin, granulocyte colony stimulating factor protein and interferon alpha
                                                                                                                                                                                                             ö
                                                        of
encoding the TR2 Orphan receptor, EAR3/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropotetin, granulocyte colony stimulating factor protein and interferon alpha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin.
                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ribozyme; erythropoietin; granulocyte colony stimulating factor; interferon alpha; ss.
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                                                                                                                                                                    Query Match 3.0%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 3.2e+02; Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17 BP; 0 A; 10 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                  Sequence 17 BP; 1 A; 8 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Blatt L, Zwick M, Pavco P, Mcswiggen J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hammerhead ribozyme substrate #1181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 37; Page 82; 164pp; English.
                                                                                                                                                                                                                                                                                                                                                                    AAF02886/c
ID AAF02886 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                    345 CGGCTGCTCTACAGCG 360
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                                                                                                                                                                                                                                                                            (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-647423/62.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens,
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Lockhart

99US-0126473P. 99US-0140359P.

RESULT

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Human; 8s; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme;
The present invention relates to an oligonucleotide array comprising oligonucleotide tags fixed to a solid substrate. The oligonucleotide array is useful for genotyping a nucleic acid sample at one or more loci via single base extension (SBE) reactions. A pair of primers is used to amplify a polymorphic locus in a sample e.g. a single nucleotide polymorphism (SNP). The present sequence is one of the primers used in the method of the present invention to amplify a polymorphic sample. The amplified nucleic acid product is then used as a template in a SBE reaction with an extension primer. The SBE reaction products are used to
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                                                                                                                                                                                                                                                                                                                                                              3.0%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 3.2e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                            Sequence 17 BP; 3 A; 7 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                              form the oligonucleotide array
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP
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Best Local Similarity 87.5
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human NOGO Inozyme #110.
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ID ABK0
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AC ABK0
XX
DT 12-M
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DE Huma
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Huma
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Huma
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KW Gere
KW musc
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Gaps

Query Match
3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels

266 GCACCTGGAGCAGGGC 281

8

GCACCGGGAGCGGGGC 1

16

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inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; Parkinson's disease; ataxia; Huntington's disease; Creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease. DNAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; B-cell Jymphoma; non-Hodgkin's lymphomy; lymphocytic leukaemia; human immunodeficiency virus; HIV associated NHL; mantle-cell lymphoma; MCL; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia;

Homo sapiens. Synthetic.

WO200159103-A2.

16-AUG-2001

09-FEB-2001; 2001WO-US004273

11-FEB-2000; 2000US-0181797P. 28-FEB-2000; 2000US-0185516P. 06-MAR-2000; 2000US-0187128P.

RIBOZYME PHARM INC. BLATT L.

(RIBO-) RIBOZYME PHARM (BLAT) BLATT L. (MCSW/) MCSWIGGEN J. (CHOW/) CHOWRIRA B M.

Chowrira BM Blatt L, Mcswiggen J,

WPI; 2001-607195/69.

Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury.

Claim 88; Page 79; 200pp; English.

The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NG20). The regulates expression of a neurite growth inhibitor gene (NG20). The CD20 mucleic acids acids eaving a an RNA molecule possessing an Indoxyme or a DNAzyme or an amberzyme (cleaving RNA with a NRN moleif) proposessing an NCH motif), a G-cleaver (cleaving RNA with a NRN moleif) proposessing an NCH motif), a G-cleaver (cleaving RNA with a NCH motif) proposessing an NCH motif) a G-cleaver (cleaving RNA with a NCH motif) proposessing an NCH motif) a gradual propose (cleaving RNA with a YG2 motif). The CD20-targetting nucleic acid is used to cleave RNA of CC of CD20 in the presence of a divalent cation that is preferably MG²+.

Furthermore, it may be contacted with a coll to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CC CD20. The treatment may further comprise the use of fone or more characterist in particular, be-call lymphoma, low-grade or follicular NHL, lymphocytic lymphoma, leukaemia, and inflammatory arthropathy. The NOGO-targetting nucleic acid may be contacted with a cell to reduce NOGO gene in the presence of a divalent cation that is preferably MG²+. Furthermore, the conclain and treat a patient having a condition associated MHL, mance of the contacted with a cell to reduce NOGO activity of the CC CC therapies. In particular, the NOGO-targetting nucleic acid may be contacted with a cell to reduce acid may be used to the captows system (NNS) injury model cation associated with the level of the contacted neuropathy, amyotrophic lateral acid caid may be contacted with a cell to reduce acid may be contacted with a cell to reduce NOGO activity of the NOGO-the treatment may further compise the use of one or more cell and treat a patient having a condition associated may be used to treat central nervous system (NNS) injury model cation accided acided or a sequence is an inozyme of the invention

Seguence 17 BP; 1 A; 8 C; 8 G; 0 T; 0 U; 0 Other;

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                                                                                                                                                                                                                         Human; ss; antisense therapy; cytostatic; antiinflammatory, haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOG0; hammerhead ribozyme; DNAzyme; inozyme; declarer; amberzyme; zinzyme; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphomy; leukaemia; human immunodeficiency virus; HTV associated NHL; mantle-cell lymphoma; MC1; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; Parkinson's disease; ataxia; Huntington's disease; creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.
                       ;
0
Length 17;
3.0%; Score 12.8; DB 1;
87.5%; Pred. No. 3.2e+02;
tive 0; Mismatches 2;
                                               302 cereaccececane 317
                                                                  2 CCGGCCCCCGGGAC 17
                                                                                                                                ABK02394 standard; RNA; 17
                                                                                                                                                                               12-MAR-2002 (first entry)
Query Match
Best Local Similarity 87.5
Matches 14; Conservative
                                                                                                                                                                                                      Human NOGO Amberzyme #66.
                                                                                                                                                       ABK02394;
                                                                                                         RESULT 453
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Synthetic.

Homo sapiens.

WO200159103-A2.

16-AUG-2001.

09-FEB-2001; 2001WO-US004273.

11-FEB-2000; 2000US-0181797P. 28-FEB-2000; 2000US-0185516P. 06-MAR-2000; 2000US-0187128P.

(RIBO-) RIBOZYME PHARM INC. (BLAT/) BLATT L. (MCSW/) MCSWIGGEN J. (CHOW/) CHOWRIRA B M.

Blatt L, Mcswiggen J, Chowrira BM;

WPI; 2001-607195/69.

Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gane or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury.

Claim 88; Page 131; 200pp; English.

The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NOGO). The nucleic acids (e.g. a ribozyme or a DNAZYME) an Inozyme (an endolytic nucleic acids (e.g. a ribozyme or a DNAZYME) an Inozyme (an endolytic nucleic acid cleaving an RNA molecule possessing an NCH mocif), a G-cleaver (cleaving RNA with a NYM mocif) pr an amberzyme (cleaving RNA with a NGM triplet), a zinzyme (cleaving RNA with a YGY motif). The CD20-atty and nucleic acid is used to cleave RNA of CD20 in the presence of a divalent cation that is preferably Mg^2^+. Purthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more

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treat lymphoma, leukaemia, be-cell lymphoma, low-grade or follicular non-treat lymphoma, leukaemia, be-cell lymphoma, low-grade or follicular non-thody in the light of the li
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Sequence 17 BP; 1 A; 9 C; 7 G; 0 T; 0 U; 0 Other;

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Gaps
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0
Match
Local Similarity 87.5%; Pred. No. 3.2e+02;
es 14; Conservative 0; Mismatches 2; Indels
                                                             305 GAGCCCCGGGGACCGC 320
      Query Match
                    Best Loca
Matches
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acacccccidedacccc 17

ABK01169 standard; RNA; 17 BP 12-MAR-2002 (first entry) Human NOGO Inozyme #439. ABK01169; RESULT 454 ABK01169,

Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NG20; hammerhead ribozyme; DNAzyme; inozyme; deleaver; amberzyme; zinzyme; linozyme; deleaver; amberzyme; linzyme; lymphoma; leukaemia; human immunodeficiency virus; HIV associated MHi, mantle-cell lymphoma; MCI; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; ampotrophic lateral sclerosis; Parkinson's disease; ataxia; Huntingcon's disease; creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease. 

sapiens Synthetic Ношо

40200159103-A2.

6-AUG-2001.

39-FEB-2001; 2001WO-US004273.

11-FEB-2000; 2000US-0181797P. 28-FEB-2000; 2000US-0185516P. 06-MAR-2000; 2000US-0187128P.

RIBOZYME PHARM INC. RIBO-> (BLAT/)

BLATT L. MCSWIGGEN J. CHOWRIRA B M. CHOM/)

Chowrira BM; Mcswiggen J, Blatt L,

WPI; 2001-607195/69. 

Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury.

Claim 88; Page 85; 200pp; English.

The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NGOD). The regulates expression of a neurite growth inhibitor gene (NGOD). The nucleic acids (e.g. a ribozyme or a nucleic acids (e.g. a ribozyme or a nucleic acids (e.g. a ribozyme or a nucleic and in nozyme (an endolytic nucleic acid cleaving an RNM molfl) proposessing an NCH molfl). The CD20-targetting RNM with a NCH will propose (cleaving RNM with a NCH will) a respective (cleaving RNM with a NCH molfl). The CD20-targetting nucleic acid is used to cleave RNA of the presence of a divalent cation that is preferably Mg² +. Furthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more therapies. In particular, the CD20 targetting nucleic acid may be used to treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular nontreavament, immunocytoma (IMC), small B-cell lymphocytic lymphoma (MCL), immunocytoma (IMC), small B-cell lymphocytic lymphoma, compania, and inflammatory arthropathy. The NOGO gene in the presence of a divalent cation that is preferably Mg² +. Furthermore, the creatment may further comprise the use of one or more cell and treat a patient having a condition associated with the level of the cell and treat a patient having a condition associated with the level of the stroke), Alzheimer's disease, dementia, multiple sclerosis (MS), chemchbrapy-induced neuropathy, and/Or other neurodegenerative disease taxia, Huntington's disease, central response of the invention of NOGO expression. The present secure is an inozyme of the invention of NOGO expression. The present sequence is an inozyme of the invention

Sequence 17 BP; 2 A; 6 C; 2 G; 0 T; 7 U; 0 Other;

Gaps ö Query Match
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels

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286 CCAAGCTGGTGAAGGA 301 쉱

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(first entry) Human NOGO Inozyme #112. 12-MAR-2002 ABK00842;

ABK00842 standard; RNA; 17 BP.

Human; 88; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; DNAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; human immunodeficiency virus; Hy associated NHL; mantle-cell lymphoma; NCL; immunocytoma; INC; immunocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; RESULT 455
ABK00842
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AC ABK0084
DT 12-MARXX
XX
HUMAN 10
KW W CSTEDFO
KW CSTEDFO
KW MASCUL
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ВЪ.

ABK02395 standard; RNA; 17

(first entry)

12-MAR-2002

ABK02395;

Human NOGO Amberzyme #67.

sapiens

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Parkinson's disease, ataxia, Huntington's disease;
Creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.
                                                              11-FEB-2000; 2000US-0181797P.
28-FEB-2000; 2000US-0185516P.
06-MAR-2000; 2000US-0187128P.
                                                    09-FEB-2001; 2001WO-US004273
                                                                                   (RIBO-) RIBOZYME PHARM INC. (BLAT/) BLATT L.
                                                                                                             Blatt L, Mcswiggen J,
                                                                                         (BLAT/) BLATT L.
(MCSW/) MCSWIGGEN J.
(CHOW/) CHOWRIRA B M.
                                                                                                                       WPI; 2001-607195/69.
                               WO200159103-A2.
                                          16-AUG-2001
                      Synthetic.
                Ношо
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Chowrira BM;

The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NG20). The regulates expression of a neurite growth inhibitor gene (NG20). The consists of an individual part of a ribozyme or a nucleic acids (e.g. a ribozyme or a nucleic acids an INCH moutle; present on the nucleic acid cleaving RNA with an NRN motif) prossessing an NCH motif), a G-cleaver (cleaving RNA with an NRN motif) prossessing an NCH motif), a G-cleaver (cleaving RNA with a NRN motif) prossessing an NCH motif), a G-cleaver (cleaving RNA with a Presence of a divalent cation that is preferably MG²+. Creat User of CD20 in the presence of a divalent cation that is preferably MG²+. Furthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of fone or more therapies. In particular, the CD20 targetting nucleic acid may be used to treat lymphoma (NCL), immunocytoma (NCL), small B-cell lymphocytic lymphoma, CCC (mucleic acid may be contacted with a cell to reduce NCGO activity of the creatment may further comprise the use of one or more presence of a divalent cation that is preferably MG²+. Furthermore, the creatment may further comprise the use of one or more cell and treat a patient having a condition associated with the level of the contacted with a cell to reduce NCGO activity of the contacted with a cell to reduce NCGO activity of the contacted with a cell to reduce NCGO activity of the contacted with a cell to reduce NCGO activity of the contacted with a cell to reduce NCGO activity of the contacted with a cell to reduce NCGO activity of the contacted with a cell to reduce NCGO activity of the contacted with a cell to reduce NCGO activity of the contacted with a cell to reduce NCGO activity of the contacted with a cell to reduce NCGO activity of the contacted neuropsethy, a Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and sequence is an inozyme of the invention Claim 88; Page 79; 200pp; English. central nervous system injury.

Sequence 17 BP; 1 A; 9 C; 7 G; 0 T; 0 U; 0 Other;

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Query Match 3.0%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 3.2e+02; Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                     303 CTGAGCCCCGGGGACC 318
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Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor sene; NOGO; hammerhead ribozyme; DNAzyme; inozyme; decleaver; amberzyme; zinzyme; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphoma; leukaemia; human immunodeficiency virus; HTV associated NHL; mantle-cell lymphoma; MCL; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; parkinson's disease; ataxia; Huntington's disease; creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chowrira BM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 88; Page 131; 200pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                     11-FEB-2000; 2000US-0181797P.
28-FEB-2000; 2000US-0185516P.
06-MAR-2000; 2000US-0187128P.
                                                                                                                                                                                                                                                                                                                                                                             09-FEB-2001; 2001WO-US004273.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            (RIBO-) RIBOZYME PHARM INC. (BLAT/) BLATT L. (MCSW/) MCSWIGGEN J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Blatt L, Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CHOWRIRA B M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-607195/69.
                                                                                                                                                                                                                                                                                                                        WO200159103-A2.
                                                                                                                                                                                                                                                                                   sapiens.
                                                                                                                                                                                                                                                                                                                                                    16-AUG-2001.
                                                                                                                                                                                                                                                                                                Synthetic.
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The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NGGO). The nucleic acids may be enzymatic nucleic acids aribozyme or a concleic acids may be enzymatic nucleic acid cleaving an RNA molecule DNAzyme) an Inozyme (an endolytic nucleic acid cleaving an RNA molecule con amberzyme (cleaving RNA acid cleaving RNA to make an amberzyme (cleaving RNA to contact an amberzyme (cleaving RNA to contact an amberzyme (cleaving RNA to contact and the acid is used to cleave RNA with a VYZ motif). The CD20-targetting nucleic acid is used to cleave RNA coff contact a patient having a condition associated with the level coff cD20. The treatment may further comprise the use of one or more contract lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular non-crow treat lymphoma (NHL), bulky low-grade or follicular non-cleukaemia, HIV (human immunodeficiency virus) associated NHL, lymphoma, contact lymphoma (NML), immunocytoma (MC), small B-cell lymphoma, contact lymphoma, contact acid massociated NHL, mantle-cell contact the contact lymphoma in the interpathy. The NOGO gene in the targetting nucleic acid is used to cleave RNA of the NOGO gene in the

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presence of a divalent cation that is preferably Mg^2^+. Furthermore, the nucleic acid may be contacted with a cell to reduce NOGO activity of the cell and treat a patient having a condition associated with the level of NOGO. The treatment may further comprise the use of one or more treat central nervous system (CMS) injury and cerebrovascular accident (CVA, stroke), Alzheimer, MSN injury and cerebrovascular accident (CVA, stroke), Alzheimer, administration accident parkinson's disease, dementia, multiple sclerosis (ALS), chemotherapy-induced neuropathy, amyocrophic lateral sclerosis (ALS), parkinson's disease, creuzfeldt-Jakob disease, muscular dystrophy, and/or other neurodegenerative disease states which respond to the modulation of NOGO expression. The present sequence is an amberzyme molecule of the invention

Sequence 17 BP; 1 A; 9 C; 6 G; 0 T; 1 U; 0 Other;

2; Indels 0; Gaps Query Match
3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels

305 GAGCCCCGGGGACCGC 320 ò g

ABN07567 RESULT

ABN07567 standard; DNA; 17 BP 

ABN07567;

29-MAY-2002 (first entry)

Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:7559.

Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.

Homo sapiens.

WO200192524-A2.

06-DEC-2001.

25-MAY-2001; 2001WO-US016981

21. SEP-2000; 2000US-0234687P.
27-SEP-2000; 2000US-0234687P.
27-SEP-2000; 2000US-0234687P.
30-JAN-2001; 2001WO-US000661.
30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 201WO-US0006663.
30-JAN-2001; 201WO-US000666.
30-JAN-2001; 201WO-US000666.
30-JAN-2001; 201WO-US000666.
30-JAN-2001; 201WO-US000666.
30-JAN-2001; 201WO-US000666.
30-JAN-2001; 201WO-US000668.

(AEOM-) AEOMICA INC.

Shannon ME Chen W, Rank DR, Hanzel DK, Gu Y, Ji Y, Penn SG, WPI; 2002-179446/23. New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.

Disclosure; SEQ ID NO 7559; 214pp; English

The present invention describes a human genome-derived myosin-like protein 1 (hgbMLP-1). The protein and polymucleotide sequences of hgbMLP-1 can be used in gene therapy and vaccine production. The hgbMLP-1 mucleic acids in samples, as amplification substrates, to nucleic acids in samples, as amplification substrates, to hgbMLP-1 mucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hgbMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hgbMLP-1 proteins or polypeptides may be concentration and/or amount specifically of hgbMLP proteins or specific blomolecule and/or amount specifically of hgbMLP proteins, as specific blomolecule capture probes for surface-enhanced laser desoxiption indisation, as therapeutic supplement in patients having specific deficiency in hgbMLP-1 production, and in vaccines or for xeplacement therapy. The production, and in vaccines or for xeplacement therapy. The production, and in vaccines or for xeplacement therapy. The production, and in vaccines or for xeplacement therapy. The production, and in vaccines or for xeplacement therapy. The production, and in vaccines or for xeplacement therapy. The production, and in vaccines and holds.

CC polymuclectide sequence expressents an oligomer used in the screening of the holds. He sequence in the exemplification of the present invention. N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from MIPO ct figure in the very production of the present invention. ô Gaps ö Query Match
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels Sequence 17 BP; 6 A; 3 C; 6 G; 2 T; 0 U; 0 Other; ABNO6000 standard; DNA; 17 BP. 385 ACGACGCCCCAAGAA 400 2 ATGACGGGCCAAGAA 17 ABN06000; RESULT 458 ABNO6000/ ઠ

Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss. Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:5992. 26-MAY-2000; 2000US-0207456P.
21-SEP-2000; 2000US-0234667P.
27-SEP-2000; 2000US-0235359P.
04-0CT-2000; 2000US-00264283.
30-JAN-2001; 2001W0-US000661.
30-JAN-2001; 2001W0-US000663.
30-JAN-2001; 2001W0-US0006663.
30-JAN-2001; 2001W0-US0006663.
30-JAN-2001; 2001W0-US0006665. 2001WO-US000668. 2001WO-US000669. 2001WO-US000670. 2001US-0266860P. 25-MAY-2001; 2001WO-US016981 29-MAY-2002 (first entry) WO200192524-A2. 30-JAN-2001; 30-JAN-2001; 05-FEB-2001; Homo sapiens. 06-DEC-2001. 

Shannon ME;

Chen W,

Hanzel DK, Rank DR,

Ji Y, Penn SG,

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30-JAN-2001; 2001WO-US000662. 30-JAN-2001; 2001WO-US000663. 30-JAN-2001; 2001WO-US000664. 30-JAN-2001; 2001WO-US000666. 30-JAN-2001; 2001WO-US000666. 30-JAN-2001; 2001WO-US000667. 30-JAN-2001; 2001WO-US000669. 30-JAN-2001; 2001WO-US000669.

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New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser descrption ionization, comprises human myosin-like protein hGDMLP-1.
(AEOM-) AEOMICA INC.
                        WPI; 2002-179446/23.
            Gu Y,
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The present invention describes a human genome-derived myosin-like

protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1

c can be used in gene therapy and vaccine production. The hGDMLP-1

nucleic acids can be used as probes to detect, characterise and quantify

c hGDMLP-1 nucleic acids in samples, as amplification substrates, to

provide initial substrates for the recombinant engineering of hGDMLP-1

protein variants having desired phenotypic improvements, and for

expressing the proteins. The hGDMLP-1 proteins or polypeptides may be

c synthesing the proteins as standards in assays used to determine the concentration

candor amount specifically of hGDMLP- proteins, as specific biomolecule

and/or amount specifically of hGDMLP proteins, as specific biomolecule

capture probes for surface-enhanced laser description ionisation, as

therapeutic supplement in patients having specific deficiency in hGDMLP-1

production, and in vaccines or for replacement therapy. The

production, and in vaccines or for replacement therapy. The

colynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a

cc disorder associated with the expression of hGDMLP-1, in particular heart

cand skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22.

The present sequence in the expression of the present invention. N.B.

CT herapeuence data for this patent did not form part of the printed

specification, but was obtained in electronic format directly from WIPO

cycly the present sequence are presented and electronic format directly from WIPO
Disclosure; SEQ ID NO 5992; 214pp; English
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## Query Match 3.0%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 3.2e+02; Matches 14; Conservative 0; Mismatches 2; Indels 351 CTCTACAGGGACTTCC 366

Sequence 17 BP; 5 A; 2 C; 7 G; 3 T; 0 U; 0 Other;

Gaps

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Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
                                                                                                                         Human GDWLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:5988.
                                                             ABN05996 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                       25-MAY-2001; 2001WO-US016981
16 CTCTACATGGACTTCC 1
                                                                                                       29-MAY-2002 (first entry)
                                                                                                                                                                                                               WO200192524-A2.
                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                   06-DEC-2001.
                                                                                 ABN05996;
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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polymucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 can be used as probes to detect, characterise and quantify mucleic acids can be used as probes to detect, characterise and quantify controlled initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for capturessing the proteins. The hGDMLP-1 proteins or polypeptides may be concentration and for amount specifically of hGDMLP-1 proteins, as specifically recognise hGDMLP-1 proteins, as specific blomolecule and/or amount specifically of hGDMLP-1 proteins, as specific blomolecule capture probes for surface-enhanced laser desorption ionisation, as capture probes for surface-enhanced laser desorption ionisation, as production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The concentration and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The concentration and in vaccines or for replacement therapy. The concentration and in vaccines or for replacement therapy. The concentration and skeleral muscle disorders, hGDMLP-1 may be used for diagnosing a confusion of the present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence data for this patent did not form part of the printed capture of the present invention. N.B. Concentration, but was obtained in electronic format directly from MIPO capture for the present directly from MIPO concentration and production and 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                                                                                                               Shannon ME;
                                                                                                                                                                                                                                                                                                                                                                                  Chen W,
                                                                                                                                                                                                                                                                                                                                                                               Hanzel DK, Rank DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; SEQ ID NO 5988; 214pp; English
                                                                                                                                                                                                                                                                                                                                                                                  Gu Y, Ji Y, Penn SG,
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                                                                                                         0; Gaps
                                                Query Match
3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
Sequence 17 BP; 6 A; 2 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                   354 TACAGCGACTTCCTCA 369
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Human GDMLP-1 17-mer scanning SEQ ID NO:4 sequence SEQ ID NO:1009. B 17 TACATGGACTTCCTCA 2 ABN01017 standard; DNA; 17 (first entry) 29-MAY-2002 ABN01017; RESULT 460 ABN01017 ઠ 셤

Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.

Homo sapiens

26-MAY-2000; 2000US-0207456P. 21-SEP-2000; 2000US-023468PP. 27-SEP-2000; 2000US-02359P. 04-0CT-2000; 2000GB-00024263. 30-JAN-2001; 2001WO-US000661.

WO200192524-A2.

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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 claim be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids in samples, as amplification substrates to the protein earlies acids in samples, as amplification substrates, to hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypoptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specifically recognise hGDMLP-1 proteins, as specific biomolecule concentration and/or amount specifically of hGDMLP proteins, as specific deficiency in hGDMLP-1 proteins, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The protein and sealences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1 is particular heart and skaletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence data for this patent did not form part of the printed sequence data for this patent did not form part of the printed sequence in the exemplification of the present invention. N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at fip., when the present and patent did not form part of the present sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chen W, Shannon ME;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17 BP; 8 A; 3 C; 6 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gu Y, Ji Y, Penn SG, Hanzel DK, Rank DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure, SEQ ID NO 1009; 214pp; English.
                                                                                                                                                                                                        30-JAN-2001; 2001WO-US00661.

30-JAN-2001; 2001WO-US006662.

30-JAN-2001; 2001WO-US000663.

30-JAN-2001; 2001WO-US000664.

30-JAN-2001; 2001WO-US000666.

30-JAN-2001; 2001WO-US000666.

30-JAN-2001; 2001WO-US000666.

30-JAN-2001; 2001WO-US000669.

30-JAN-2001; 2001WO-US000669.

30-JAN-2001; 2001WO-US000669.
                                                                                                                                              2000US-0234687P,
2000US-0236359P,
2000GB-00024263,
                                                                               25-MAY-2001; 2001WO-US016981
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (AEOM-) AEOMICA INC.
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                                    06-DEC-2001
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Gaps
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Query Match 3.0%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 3.2e+02; Matches 14; Conservative 0; Mismatches 2; Indels
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ABN01018 standard; DNA; 17 BP. ABN01018; RESULT 461
ABNO1018
ID ABNO103
XX
AC ABNO103

Human GDMLP-1 17-mer scanning SEQ ID NO:4 sequence SEQ ID NO:1010. (first entry) 29-MAY-2002

Human, genome-derived myosin-like protein 1, GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.

Homo sapiens.

40200192524-A2

25-MAY-2001; 2001WO-US016981

26-MAY-2000;

27-58P-2000; 2000US-0236359F 04-0CT-2000; 2000GB-0236359F 30-JAN-2001; 2001W0-US000661. 30-JAN-2001; 2001W0-US000661. 30-JAN-2001; 2001W0-US000663. 30-JAN-2001; 2001W0-US0006664. 30-JAN-2001; 2001W0-US0006665. 30-JAN-2001; 2001W0-US0006667. 30-JAN-2001; 2001W0-US0006669. 30-JAN-2001; 2001W0-US0006669.

05-FEB-2001; 2001US-0266860P.

(AEOM-) AEOMICA INC.

Shannon ME; Gu Y, Ji Y, Penn SG, Hanzel DK, Rank DR, Chen W, WPI; 2002-179446/23. New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.

Disclosure, SEQ ID NO 1010; 214pp; English.

The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polymucleotide sequences of hGDMLP-1 can be used as probes to detect, characterise and quantify and vaccine production. The hGDMLP-1 nucleic acids in samples, as amplification substrates, to hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as the represent in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production and with the expression of hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1 is localised to chromosome 22.

The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at fits wipo.int/pub/published_pot_esquence

Sequence 17 BP; 8 A; 2 C; 7 G; 0 T; 0 U; 0 Other;

Gaps ö Query Match
3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels

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wed Apr 41 14:56:41 400%
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ABN07571 standard; DNA; 17 BP 29-MAY-2002 (first entry) ABN07571; RESULT 

Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:7563.

Human, genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.

Homo sapiens

WO200192524-A2.

06-DEC-2001.

25-MAY-2001; 2001WO-US016981

26-MAY-2000; 2000US-0207456P.
21-SEP-2000; 2000US-0234637P.
27-SEP-2000; 2000US-0234535P.
04-OCT-2000; 2000US-00024263.
30-JAN-2001; 2001WO-US000661.
30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000667.

(AEOM-) AEOMICA INC.

DK, Rank DR, Chen W, Shannon ME; Hanzel Gu Y, Ji Y, Penn SG, WPI; 2002-179446/23. New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.

Disclosure; SEQ ID NO 7563; 214pp; English.

The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1): The protein and polymucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 mucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 mucleic acids in samples, as amplification substrates to hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be cased as immunospens for raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser described biomolecule therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production and in vaccines of for replacement therapy. The polymolectide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart can disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22.

The present sequence represents an oligomer used in the screening of the hGDWLP-1 sequence in the exemplification of the present invantion. N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequence 8888888

Sequence 17 BP; 6 A; 3 C; 7 G; 1 T; 0 U; 0 Other;

ö ; 0 3.0%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 3.2e+02; 7ative 0; Mismatches 2; Indels Best Local Similarity 87.5 Matches 14; Conservative Query Match

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ABK26660 standard; DNA; 17 BP.

ABK26660;

09-APR-2002 (first entry)

Waxy starch production genome altering oligonucleotide #316.

Chromosomal genomic alteration, genome altering oligonucleotide; PCR; ss; o-methyl modification, LNA modification; phosphorothioate linkage; DNA repair; DNA alteration; environmental tolerance; hygromycinh. Brimer; abiotic stress tolerance; improved nutritional value; hygromycinh. Brimer; amino acid over production; herbicide resistance; hygromycin, hygromycin; porphyric herbicide resistance; sulphonylurea herbicide resistance; porphyric herbicide resistance; sulphonylurea herbicide resistance; modified oil production; modified starch production; waxy starch; altered floral morphology; male-sterile plant; albino mutant; andified fatty acid content; reduced palmitate production; albino plant; increased stearate production; reduced linolenic acid production; ABRULT 46
ABRULT 46
ABRULT ABR

Oryza sativa. Synthetic. WO200192512-A2.

06-DEC-2001.

01-JUN-2001; 2001WO-US017672.

01-JUN-2000; 2000US-0208538P. 30-OCT-2000; 2000US-0244989P. 27-MAR-2001; 2001US-00818875.

(UYDE ) UNIV DELAWARE.

Rice MC, Kim Kmiec EB, Gamper HB,

WPI; 2002-106307/14.

New oligonucleotides with modified nuclease-resistant termini, useful for creating plants with desired phenotypes, e.g. stress tolerance, improved nutritional value, herbicide or disease resistance, or modified oil production.

Claim 7; Page 163; 220pp; English.

The invention relates to an oligonucleotide for targeted alteration of a genetic sequence, which comprises a single-stranded oligonucleotide having a DNA domain. The DNA domain has at least one mismatch with respect to the genetic sequence to be altered and further comprises chemical modifications of the oligonucleotide. The chemical modifications consist of o-methyl modification, an LNA modification, two or more phosphorothioate linkages on a terminus, or a combination of any two or

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directing repair or alterations. The oligonucleotides are useful for directing repair or alteration of plant genetic information. The coligonucleotides are particularly useful for recating plants with desired phenotypes, e.g. environmental or abilotic stress tolerance, improved contritional value (e.g. altering amino acid content of plants or conferring amino acid over production), herbicide resistance (e.g. offersting amino acid over production), herbicide resistance (e.g. g. g. glyphosate resistance, imidazolinone and sulphonylurea herbicide casistance), classase resistance, nonfified resistance or triazine resistance), disease resistance, modified atarch production of waxy starch), altered floral corphology (e.g. male-sterile plants) or modified fatty acid content corphology (e.g. male-sterile plants) or modified fatty acid content ceg. increased palmitate, increased stearate or reduced linolenic acid). The oligonucleotides are also useful for producing albino mutants for the analysis of photosynthetic processes. This sequence represents a genome content content canalysis of photosynthetic processes. This sequence represents a genome content canalysis of photosynthetic processes. This sequence represents a genome content canalysis of photosynthetic processes. This sequence represents a genome content canalysis of photosynthetic processes.

Sequence 17 BP; 3 A; 7 C; 6 G; 1 T; 0 U; 0 Other;

Gaps ; 3.0%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 3.2e+02; ative 0; Mismatches 2; Indels Local Similarity 87.5 nes 14, Conservative Query Match Matches

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2 cadcdacracddddd 17

5639/c ABK26639 standard; DNA; 17 BP. ABK26639; ABK26639, RESULT THE LEAD TO BE SEEN THE CONTRACT OF THE PROPERTY OF THE PROPER

Waxy starch production genome altering oligonucleotide #295.

Chromosomal genomic alteration; genome altering oligonucleotide; PCR; ss; o-methyl modification; LNA modification; phosphorothioate linkage; DNA regair; DNA alteration; environmental tolerance; hygromycin; abiotic stress tolerance; improved mutritional value; hygromycin; primer; amino acid over production; herbicide resistance; glyphosate resistance; infazzolinone herbicide resistance; sulphonylurea herbicide resistance; porphyric herbicide resistance; triazine resistance; disease resistance; altered floral modified starch production; waxy starch; altered floral morphology; male-sterile plant; albino mutant; modified fatty acid content; reduced plant; albino mutant; increased stearate production; reduced linolenic acid production; albino plant; increased stearate production; reduced linolenic acid production; photosynthetic process.

Synthetic

01-JUN-2001; 2001WO-US017672

01-JUN-2000; 2000US-0208538P. 30-OCT-2000; 2000US-0244989P. 27-MAR-2001; 2001US-00818875.

Kim J; Kmiec EB, Gamper HB, Rice MC, New oligonucleotides with modified nuclease-resistant termini, useful for creating plants with desired phenotypes, e.g. stress tolerance, improved nutritional value, herbicide or disease resistance, or modified oil production.

Claim 7; Page 162; 220pp; English.

The invention relates to an oligomuclectide for targeted alteration of a genetic sequence, which comprises a single-stranded oligomuclectide comparises a single-stranded oligomuclectide having a DNA domain has at least one mismatch with respect to the genetic sequence to be altered and further comprises consists of o-methyl modification, an INA modification, two or more consists of o-methyl modification, an INA modification two or more phosphorothicate linkages on a terminus, or a combination of any two or more of these modifications. The oligomuclectides are useful for configuration repair or alteration of plant genetic information. The oligomuclectides are particularly useful for creating plants with desired of phenotypes, e.g. environmental or abiotic stress tolerance, improved conferring amino acid over production), herbicide resistance, indazolinone and sulphonylurea herbicide conferring amino acid over production, herbicide resistance, indazolinone and sulphonylurea herbicide conferring and starch or production, modified fatry acid content (e.g. increased starch or production, modified fatry acid content (c.g. increased starch or production of waxy starch), altered floral morphology (e.g. male-sterile plants) or modified fatry acid content (c.g. increased starch increased stearts or reduced linolenic acid). The oligomuclectides are alse ouseful for production mutants for the analysis of photosynthetic processes. This sequence represents a genome content of the invention content of the inventical content of the inve

Sequence 17 BP; 1 A; 6 C; 7 G; 3 T; 0 U; 0 Other;

.; 0 3.0%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 3.2e+02; Live 0; Mismatches 2; Indels Query Match Best Local Similarity 87.5 Matches 14; Conservative

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8 셤 ABK26659 standard; DNA; 17 BP.

ABK26659;

09-APR-2002 (first entry)

Waxy starch production genome altering oligonucleotide #315.

Chromosomal genomic alteration; genome altering oligonucleotide; PCR; ss; o-methyl modification; LNA modification; phosphorothioate linkage; buby repairs in the alteration; environmental tolerance; hygromycin. Bit abicic stress tolerance; improved mutritional value; hygromycin. Brimary. Indiance herbicide resistance; glyphosate resistance; porphyric herbicide resistance; sulphnorilurea herbicide resistance; porphyric herbicide resistance; triazine resistance; modified oil production; modified starch production; waxy starch; modified floral morphology; male-sterile plant; albino mutant; modified attach production; abbino mutant; increased stearate production; reduced palmitate production; albino plant; increased stearate production; reduced linolenic acid production; photosynthetic process. 

Oryza sativa. Synthetic.

WO200192512-A2.

01-JUN-2001; 2001WO-US017672. 06-DEC-2001.

2000US-0208538P. 2000US-0244989P. 2001US-00818875. 01-JUN-2000; 2 30-OCT-2000; 2 27-MAR-2001; 2

09-APR-2002 (first entry)

Oryza glaberrima.

WO200192512-A2.

06-DEC-2001

(UYDE ) UNIV DELAWARE.

WPI; 2002-106307/14.

380 CCGCGACGACGCGCC 395

ठे g

Kim J;

Rice MC,

New oligonucleotides with modified nuclease-resistant termini, useful for creating plants with desired phenotypes, e.g. stress tolerance, improved nutritional value, herbicide or disease resistance, or modified oil Kmiec EB, Gamper HB, (UYDE ) UNIV DELAWARE WPI; 2002-106307/14. production. 

Claim 7; Page 163; 220pp; English.

The invention relates to an oligomuclectide for targeted alteration of a genetic sequence, which comprises a single-stranded oligomuclectide having about Amenia. The DNA domain has at least one mismatch with respect to the genetic sequence to be altered and further comprises chemical modifications of the oligomuclectide. The chemical modifications of consist of o-methyl modification, an LNA modification, two or more phosphorothicate linkages on a terminus, or a combination of any two or more of these modifications. The oligomuclectides are useful for directing repair or alteration of plant genetic information. The coligomuclectides are particularly useful for creating plants with desired phenotypes, e.g. environmental or abiotic stress tolerance, improved nutritional value (e.g. altering amino acid content of plants or conferring amino acid over production), herbicide resistance, indazolinone and sulphomylurea herbicide conferring amino acid seases resistance, indazolinone and sulphomylure production (e.g. increased starch or production), modified farty acid content (e.g. increased starch or production) of waxy starch), altered floral morphology (e.g. male-sterile plants) or modified fatty acid content (e.g. increased starch or production of waxy starch), altered floral cortent certical plants or production modified alternor analysis of photosynthetic processes. This sequence represents a genome cortent or production analysis of photosynthetic processes. This sequence represents a genome cortent or production of wax and production and production or content or content or production of wax starch), altering oligonuclectides are also useful for producting albino mutants for the altering oligonuclectides of the invention

Sequence 17 BP; 1 A; 6 C; 7 G; 3 T; 0 U; 0 Other;

Gaps ó 3.0%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 3.2e+02; rative 0; Mismatches 2; Indel8 Query Match
Best Local Similarity 87.55

ઠ a RESULT 466 ABK2664

ABK26640 standard; DNA; 17 BP. ABK26640; 09-APR-2002 (first entry)

Waxy starch production genome altering oligonucleotide #296.

Chromosomal genomic alteration; genome altering oligonucleotide; PCR; ss., o-methyl modification; LNA modification; phosphorothioate linkage; DNA repair; DNA alteration; environmental tolerance; hygromycin; abiotic stress tolerance; improved nutritional value, hygromycin; primary amino acid over production; herbicide resistance; glyphosate resistance; porphyxic herbicide resistance; sulphonylurea herbicide resistance; porphyxic herbicide resistance; sulphonylurea herbicide resistance; modified oil production; modified starch production; waxy starch; modified facty acid content; reduced plant; albino mutant; modified facty acid content; reduced plant; albino mutant; increased stearate production; reduced linolenic acid production; plant; increased stearate production; reduced linolenic acid production; photosynthetic process 

Oryza glaberrima. Synthetic.

WO200192512-A2 06-DEC-2001, 01-JUN-2001; 2001WO-US017672

01-JUN-2000; 2000US-0208538P. 30-OCT-2000; 2000US-0244989P. 27-MAR-2001; 2001US-00818875.

(UYDE ) UNIV DELAWARE.

Gamper HB, Rice MC, Kim J; Kmiec EB,

WPI; 2002-106307/14.

New oligonucleotides with modified nuclease-resistant termini, useful for creating plants with desired phenotypes, e.g. stress tolerance, improved nutritional value, herbicide or disease resistance, or modified oil production.

Claim 7; Page 162; 220pp; English.

The invention relates to an oligonuclectide for targeted alteration of a genetic sequence, which comprises a single-stranded oligonuclectide having energy and a single-stranded oligonuclectide having a DNA domain. The DNA domain has at least one mismatch with respect to the genetic sequence to be altered and further comprises chemical modifications of the oligonuclectide. The chemical modifications of consist of o-methyl modification, an LNA modification, two or more of phosphorothioate linkages on a terminus, or a combination of any two or more of these modifications. The oligonuclectides are useful for directing repair or alteration of plant genetic information. The oligonuclectides are particularly useful for creating plants with desired oligonuclecting amino acid content of plants or nutritional value (e.g. altering amino acid content of plants or conferring amino acid over production, herbicide resistance (e.g. conferring amino acid over production, herbicide resistance (e.g. cyphynosate resistance, inidazolinone and sulphomylurea herbicide resistance, or production of waxy starch), altered floral morphology (e.g. male-sterile plants) or modified fatty acid content of e.g. increased starch or production of waxy starch), altered floral corphology (e.g. male-sterile plants) or modified fatty acid content of e.g. increased starch or production of waxy starch), altered floral corphology (e.g. male-sterile plants) or modified fatty acid content of e.g. nealestance are also useful for production and anylysis of photosynthetic processes. This sequence represents a genome of altering oligonuclectide of the invention 

Sequence 17 BP; 3 A; 7 C; 6 G; 1 T; 0 U; 0 Other;

Gaps ö Query Match
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels

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380 CCGCGACGACGCGCGC 395 2 CAGCGACTACGGCGCC 17 δ 셤

ABV79109 standard; DNA; 17 BP RESULT 467 ABV79109

03-JAN-2003 (first entry) ABV79109;

Human HTPL scanning oligonucleotide SEQ ID 355.

Human, gene therapy, tumour suppressor; HTPL; chromosome 10p12.1; human testis expressed Patched like protein; testis; adrenal; liver; male germ cell development; bone marrow; brain; kidney; lung; placenta; prostate; skeletal muscle; colon; male infertility; cancer; ss. 8×4×4×4×4×4×4×

Homo sapiens

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The present invention relates to human testis expressed Patched like protein (HTPL, see ABV7875 to ABV78762 and ABB9819 to ABB9820). HTPL protein (HTPL, see ABV7875 to ABV78762 and ABB9819 to ABB98520). HTPL has two isoforms, with a few single base pair differences between the two. One of the single base pair changes introduces a premature stop codon in HTPL-8 (8 for short) compared to HTPL-16 (1 for long). HTPL shares an overall structure organisation with the Patched protein. The shares an overall structure organisation with the Patched protein. The care of that of Patched, and is a potential tumour suppressor. HTPL is not to that of Patched, and is a potential tumour suppressor. HTPL is contact in regulating male germ cell development, and the HTPL gene was important in regulating male germ cell development, and the HTPL, and in therapy and manufacture of a medicament for treatment or prevention of therapy and manufacture of a medicament for treatment or prevention of therapy and manufacture of a medicament for treatment or prevention of the HTPL. Such disorder sancdude disorders of testis, or adrenal, adult and fortal muscle or colon function. HTPL proteins and mucleic acids are clinically useful diagnostic markers and potential therapeutic agents for male infertility and cancer. The present oligonucleotide was used in an example from the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17 BP; 1 A; 8 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local ?
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0; Gaps
3.0%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 3.2e+02; tive 0; Mismatches 2; Indels
                                                     Matches 14; Conservative
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RESULT 468
ABV79107
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ABV79107 standard; DNA; 17 BP
                 03-JAN-2003 (first entry)
         ABV79107;
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Human; gene therapy; tumour suppressor; HTPL; chromosome 10p12.1; human testis expressed Patched like protein; testis; adrenal; liver; male germ cell development; bone marrow; brain; kidney; lung; placenta; prostate; skeletal muscle; colon; male infertility; cancer; ss. Human HTPL scanning oligonucleotide SEQ ID 353.

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30-JAN-2001, 2001WO-US000663.
30-JAN-2001, 2001WO-US000664.
30-JAN-2001, 2001WO-US000665.
30-JAN-2001, 2001WO-US000665.
30-JAN-2001, 2001WO-US000668.
33-JAN-2001, 2001WS-08000669.
23-MAY-2001, 2001US-00864761.
09-OCT-2001, 2001US-0327898P.
                                  28-JAN-2002; 2002EP-00001167
                                                                                          (AEOM-) AEOMICA INC
      Homo sapiens
               EP1229046-A2
                         07-AUG-2002.
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30-JAN-2001; 2001WO-US000663. 30-JAN-2001; 2001WO-US000665. 30-JAN-2001; 2001WO-US000665. 30-JAN-2001; 2001WO-US000668. 30-JAN-2001; 2001WO-US000668. 23-MAY-2001; 2001WO-US000669.

28-JAN-2002; 2002EP-00001167

EP1229046-A2 07-AUG-2002 Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL.

WPI; 2002-676582/73

Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL.

WPI; 2002-676582/73.

Zhan J;

(AEOM-) AEOMICA INC.

Example 2; Page 110; 718pp; English

Zhan J;

Example 2; Page 110; 718pp; English

The present invention relates to human testis expressed Patched like CC protein (HTPL, see ABV78759 to ABV8762 and ABB88519 to ABB88520). HTPL protein (HTPL, see ABV78759 to ABV8762 and ABB88519 to ABB88520). HTPL brotein (HTPL, see ABV78759 to ABV88612 to ABB88520). HTPL CC has two isoforms, with a few single base pair differences between the cwo. One of the single base pair changes introduces a premature stop codon in HTPL-8 (S for short) compared to HTPL-16. Lf for long). HTPL compared structural features strongly imply that HTPL plays a role similar to that of Patched, and is a potential tumour suppressor. HTPL is important in regulating male germ cell development, and the HTPL gene was important in regulating male germ cell development, and the HTPL, and in cueful for diagnosing a disorder caused by mutation in HTPL, and in therapy and manufacture of a medicament for treatment or prevention of therapy and manufacture of a medicament for treatment or prevention of therapy and manufacture of a medicament for treatment or prevention of the HTPL. Such disorder associated with decreased expression or activity of human compared or colon function. HTPL proteins and nucleic acids are shelteral muscle or colon function. HTPL proteins and nucleic acids are clinically useful diagnostic markers and potential therapeutic agents for example from the invention

0; Gaps 3.0%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 3.2e+02; Live 0; Mismatches 2; Indels Sequence 17 BP; 1 A; 9 C; 5 G; 2 T; 0 U; 0 Other; 3.0 Best Local Similarity 87.5 Matches 14, Conservative

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ABK18437 standard; RNA; 17 BP.
136 ccccccrccccccqcqq 151
             2 CCCGCCTGCCGCTGGA 17
                                                                                                                                   09-APR-2002 (first entry)
                                                                                                            ABK18437;
                                                            RESULT 469
ABK18437/c
D ABK1841
XX ABK1841
XX ABK1841
XX XX BB Human J
XX XX KX Human J
XX KW Human J
XX KW Human J
XX KW Human J
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  à
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Human; hammerhead ribozyme; cytostatic; antitumour; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; virucide; osteopathic; Human ERG hammerhead ribozyme target sequence, Seq ID No 1084.

rng.res

Novel polynucleotide which down regulates expression of Ets-related gene, useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome. vulnerary; cancer; lymphoma; Ewing's sarcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous sclerosis; port-wine stain; wound healing; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; ss; Osler-Weber-rendu syndrome, leukaemia; osteoporosis; DNAzyme; inozyme; Jarvis T, Von Carlowitz I, Mcswiggen JA, Mclaughlin F, Randi AM; 16-MAY-2001; 2001WO-US015866 16-MAY-2000; 2000US-00572021 (RIBO-) RIBOZYME PHARM INC. (GLAX ) GLAXO GROUP LTD. WPI; 2002-082995/11. WO200188124-A2 Homo sapiens. 22-NOV-2001 amberzyme.

Claim 4; Page 78; 149pp; English.

The invention relates to a nucleic acid molecule (I) which down regulates expression of an Ets-related gene (ERG). (I) is useful for treating conditions selected from cancer, lymphoma, Ewing's sarcoma, melanoma, conditions selected from cancer, lymphoma, Ewing's sarcoma, melanoma, tumour angiogenesis, diabetic retinopathy, macular degeneration, cumour angiogenesis, diabetic retinopathy, macular degeneration, conversation, arthritis, psoriasis, verruca vulgaris, angiofibroma of tubercous sclerosis, port-wine stains, Sturge weber syndrome, leukaemia, osteoprosis and wound healing. (I) is useful for treating a patient having a condition associated with the level of ERG, by contacting cells of the patient with (I) under conditions suitable for the treatment. The method comprises the use of one or more theraples to the treatment is treated by administering (I) to the patient in conjunction with one or more of other theraples such as radiation or conjunction with one or more of other theraples such as radiation or conjunction with one or more of other theraples such as radiation or conjunction with one or more of other theraples such as radiation or call, by contacting (I) is useful for reducing ERG activity in a cell, by contacting (I) with RMJ, in the presence of a divalent confiseases related to the expression of ERG, and as diagnostic tool to examine genetic drift and mutations within diseased cells or to detect the presence of ERG RNA in a cell. (I) is useful for specifically contacting genes that share homology within diseased cells or conditions and consymptic mucleic acid molecules which regulate expression of ERG, and conjunction of the invention

Sequence 17 BP; 2 A; 4 C; 6 G; 0 T; 5 U; 0 Other;

Gaps ; 0 Query Match
3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels

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RESULT 470 ABK18438/C ID ABK18438 standard; RNA; 17 BP. XC ABK18438;

Human, hammerhead ribozyme; cytostatic; antitumour; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; virucide; osteopathic; vulnerary; cancer; lymphoma; Ewing's sarcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous sclerosis; port-wine stain; wound healing; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; se; Osler-Weber-rendu syndrome; leukaemia; osteoporosis; DNAzyme; inozyme; Human ERG hammerhead ribozyme target sequence, Seg ID No 1085.

09-APR-2002 (first entry)

Homo sapiens. 

amberzyme.

WO200188124-A2.

22-NOV-2001.

16-MAY-2001; 2001WO-US015866.

16-MAY-2000; 2000US-00572021.

(RIBO-) RIBOZYME PHARM INC. (GLAX ) GLAXO GROUP LTD.

Randi AM; Mcswiggen JA, Mclaughlin F, Jarvis I, Von Carlowitz I,

WPI; 2002-082995/11.

The invention relates to a nucleic acid molecule (I) which down regulates corpression of an Eta-related gene (ERG). (I) is useful for treating capression of an Eta-related gene (ERG). (I) is useful for treating conditions selected from cancer, lymphoma, Ewing's sarcoma, melanoma, tumour angiogenesis, diabetic retinopathy, macular degeneration, cumour angiogenesis, diabetic retinopathy, macular degeneration, cumour angiogenesis, diabetic retinopathy, macular degeneration, cumour angiogenesis, versuce where syndrome, leukaemia, osteoporosis and wound healing. (I) is useful for treating a patient having a condition associated with the level of ERG, by contacting cells of the patient with (I) under conditions suitable for the treatment. The method comprises the use of one or more therapies or the treatment. Heukaemia or tumour conditions suitable for the treatment. Leukaemia or tumour conjunction with one or more of other therapies such as radiation or chaptonesis is treated by administering (I) to the patient in conjunction with one or more of other therapies such as radiation or chaptonesis is treated by administering (I) to the patient in conjunction with one or more of other therapies such as radiation or chaptonesis is treated by administering (I) to the patient in conjunction with one or more of other therapies such as radiation or chaptons with the presence of advalent cation such as Mg2+. (I) is useful for diagnosis of conditions and cation such as Mg2+. (I) is useful for diagnosis of conditions and cation such as Mg2+. (I) is useful for diagnosis of conditions or the expression of ERG fusion genes. ABK13354-ABK22719 represent mucleic acids, including antisense and cargamatic uncleic acid molecules which regulate expression of ERG, and corpuse of the invention

Sequence 17 BP; 2 A; 4 C; 5 G; 0 T; 6 U; 0 Other;

. 0 Query Match
3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels

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Gарв

16 GCCAAGAAGGCCATCT

Novel polynucleotide which down regulates expression of Ets-related gene, useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome.

Claim 4; Page 78; 149pp; English.

408 393 GCCAAGAAGGTCTTCT

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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino acids (SI, ABB83999), a sequence having 65% sequence of 6711, (S1), having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GrPases as well as downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (I) and nucleic acids (II) for identifying a specific binding partner. (I) and nucleic acids (II) are useful for adiagnoshing, monitoring disagnoshing and treating caused by altered expression of human POSHL1 including diagnoshing and treating caused by altered expression of human POSHL1 including diagnoshing are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The customer is sequence is that of a scanning oligonucleotide useful in examples of the invention. Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to berwent by the European Patent Office
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL -1, useful for treating disorders associated with decreased expression or activity of human POSHL1.
                                                                                                                                                                                                                                                                                                                                  Human; POSHL 1; SH3 domain; POSH-like signalling protein 1; oncogene;
Rho GTPase; signal transduction; gene expression; cancer; vaccine;
gene therapy; transgenic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 2; SEQ ID NO 1747; 60pp + Sequence Listing; English.
                                                                                                                                                                                                                                                       Human POSHL1 scanning oligonucleotide SEQ ID NO 1747.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17 BP; 4 A; 7 C; 5 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001US-00864761.
ABV91034 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-JAN-2002; 2002EP-00001165
                                                                                                                                                              23-DEC-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-684061/74.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (AEOM-) AEOMICA INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EP1239051-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11-SEP-2002,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Shannon M;
                                                                                    ABV91034;
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Inoko H, Kagiya T, Ichihara T, Matsumura Y, Moriya S, Nishida M;

01-JUN-2000; 2000JP-00164798.

(NISN ) NISSHINBO IND INC. (SYST-) SYSTEM RES INC.

WPI; 2002-122074/16.

01-JUN-2001; 2001WO-JP004662.

WO200192572-A1. Homo sapiens

06-DEC-2001.

Human, human leukocyte antigen; HLA; genotype; polymorphism; immunogenetic; transplantation; genetic disease; ss.

Human HLA genotyping oligonucleotide SEQ ID NO 27

21-MAR-2002 (first entry)

ABL30538;

ABL30538 standard; DNA; 17 BP

ABL30538

339 CAGGGCCGCCTGCTCT 354

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17 CAGGGCGGCTGTGCT

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The invention relates to a typing kit for judging human leukocyte antigen (HLA) genotype of a sample by hybridising a substrate on which 10-24 base oligonucleotides (ABL30512-ABL31809) originating in the sequences of genes e.g. belonging to HLA class I antigens on human genome and containing gene polymorphisms as alloantigens have been immobilised as primers for amplification of cleaved nucleic acids relating to gene polymorphisms. The method is useful for judging HLA genotypes of individuals by determining immunogenetic differences before transplanting between them, providing genetic information to decide compatibility of organ and tissue for transplantation e.g. of bone marrow, kidney, liver, pancreas, Langerhans islet in pancreas and cornea, susceptibility diagnosis of genetic diseases and identifying individuals
                                                                                                                                                                                                                                                                                                                                                                                                                                  Human leukocyte antigen (HLA) typing, useful for judging HLA genotypes of individuals e.g. by determining immunogenetic differences when transplanting between them.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          135 GCCCGCCTGGCGGTGG 150
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Gaps ö

Query Match 3.0%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 3.2e+02; Matches 14; Conservative 0; Mismatches 2; Indels

sub-unit modulating amberzyme substrate #174

Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; d-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; oesophageal cancer; stomach cancer; colorectal cancer; pancreatic cancer; oesophageal cancer; head and neck cancer; ovarian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; docetaxel; cisplatin; methotrexate; chemotherapy; paclitaxel; docetaxel; cisplatin; methotrexate; gencitabine; radarion therapy; inflammatory disease; asthma; diabetes; rheumatoid arthritis; restenosis; Crohn's disease; obesity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; seepsis; transplant/graft rejection; reperfusion inluy; glomerulonephitis; allergic airway inflammation; inflammatory bowel disease; infection; ss.

Homo sapiens.

US2002177568-A1.

28-NOV-2002

23-MAY-2001; 2001US-00864785

92US-00987132. 94US-00245466. 94US-00291932. 96US-00777916. 07-DEC-1992; 18-MAY-1994; 15-AUG-1994; 23-DEC-1996;

(STIN/) STINCHCOMB D T. (MCSW/) MCSWIGGEN J. (DRAP/) DRAPER K G.

Stinchcomb DT, Mcswiggen J, Draper KG;

WPI; 2003-340953/32.

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.

Claim 3; Page 54; 72pp; English.

The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor regulates expression of a sequence encoding a subunit of nuclear factor regulation. The enzymatic nucleic acid molecule is adapted to treat configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A.

(I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially Mg^2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or calling reastant cancer. The method involves use of other drug therapises such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, cyclophosphamide, docxoubin, fluorouracil carboplatin, edetrexate, gencitabine or radiation therapy. The enzymatic and antisense nucleic acid molecules are also useful for treating inflammatory disease such as colorestal, restenosis, asthma, crobin's disease such as colosisty, autoimmune disease, lupus, multiple solerosis, transplant/graft rejection, gene therapy applications, ischaemiar/seperfusion injury sepsis, allergic airway inflammation, inflammatory bowel disease or infection. This sequence represents the substrate of a novel enzymatic contraction.

Sequence 17 BP; 5 A; 3 C; 8 G; 0 T; 1 U; 0 Other;

ô Gaps ö 3.0%; Score 12.8; DB 1; Length 17; llarity 81.2%; Pred. No. 3.2e+02; Conservative 1; Mismatches 2; Indels Query Match Best Local Similarity Matches 13; Conserv

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286 CCAAGCTGGTGAAGGA 301
                 1 ccaggcuggggaagga
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RESULT 474 ACA06662

ACA06662 standard; RNA; 17 

ACA06662;

03-JUN-2003 (first entry)

NFKB sub-unit modulating inozyme substrate #481.

Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; broate cancer; colorectal cancer; brain cancer; cesophageal cancer; stomach cancer; bladder cancer; pancreatic cancer; cervical cancer; head and neck cancer; ovarian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; pactitaxel; docetaxel; cisplatin; methorrexate; cyclophosphanide; docetaxel; cisplatin; methorrexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatorid arthritis; restenosis; Crohn's disease; obesity; lackhemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection; ss.

Homo sapiens.

US2002177568-A1.

28-NOV-2002.

23-MAY-2001; 2001US-00864785

92US-00987132. 94US-00245466. 94US-00291932. 96US-00777916.

18-MAY-1994; 15-AUG-1994; 23-DEC-1996;

STIN/) STINCHCOMB D T. ۲, (MCSW/) MCSWIGGEN J (DRAP/) DRAPER K G.

Stinchcomb DT, Mcswiggen J, Draper KG;

WPI; 2003-340953/32.

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.

Claim 3; Page 34; 72pp; English.

rhe invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NREM), where (I) is an inozyme, zinzyme, G-Claever or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially MG-2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or cherapic such as monoclonal antibodies, REL-A-specific inhibitors or cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate,

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Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.
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acid molecules are also useful for treating inflammatory disease such as rheumatoid arthritis, restemosis, asthma, Crohm's disease, diabetes, obesity, autoimmune disease, lupus, multiple sclerosis, transplant/graft rejection, gene therapy applications, ischaemia/reperfusion injury (central nervous system (CNS) and myocardial), glomerulonephritis, espesis, allergic airway inflammation, inflammatory bowel disease or infection. This sequence represents the substrate of a novel enzymatic nucleic acid molecule
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; oesophageal cancer; stomach cancer; bladder cancer; pancreatic cancer; cervical cancer; head and neck cancer; ovarian cancer; malanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; common expressy; paclitaxal; docetaxel; cisplatin; methortexate; colophosphanide; docetaxel; cisplatin; methortexate; cyclophosphanide; docetaxel; cisplatin; detrexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatoid arthritis; restences; Crohn's disease; deshim; diabetes; rheumatoid arthritis; restences; lupus; multiple sclerosis; sechsemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sechsemia; allergic airway inflammation; inflammatory bowel disease; infection; ss.
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                                                                                                                                                                                Sequence 17 BP; 0 A; 7 C; 9 G; 0 T; 1 U; 0 Other;
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94US-00245466.
94US-00291932.
96US-00777916.
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MCSW/) MCSWIGGEN J.
DRAP/) DRAPER K G.
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15-AUG-1994;
23-DEC-1996;
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configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. treating a patient having a condition associated with the level of REL-A. The presence of advalent cation, especially MG-2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, doctaxel, displant, methotrexate, gencitabine or radiation therapy. The enzymatic and antisense nucleic cancel molecules are also useful for treating inflammatory disease such as chemotherapy as also useful for treating inflammatory disease such as the manual disease, lupus, multiple sclerosis, transplant/graft rejection, gene therapy applications, ischaemia/repeaffusion injury rejection, alergic alrway inflammatory bowel disease or sepsis, allergic alrway inflammation, inflammatory bowel disease or infection. This sequence represents the substrate of a novel enzymatic
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94US-00245466.
94US-00291932.
96US-00777916.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                nucleic acid molecule
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15-AUG-1994;
23-DEC-1996;
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ACA06661/c
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requiration. The enzymatic nucleuc acts miscure (1) which down configuration. The enzymatic nucleic acid molecule (1) which where (1) is an inozyme, zinzyme; G-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regularing REL-A activity in a cell, for treating a sequence of REL-A gene, in the presence of a divalent cation, especially MG'2+. The enzymatic and the presence of a divalent cation, especially MG'2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, corvical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies (REL-A-specific inhibitors or chemotherapy including paclitaxe), docetaxel, cisplath, methotrexate, colophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, desense nucleic acid molecules are also useful for treating inflammatory disease such as rhematoid arthritis, restenosis, asthma, Crohn's disease such as rejection, gene therapy applications, ischaemia/reperfusion injury cepterion, gene therapy applications, ischaemia/reperfusion injury cepterion, and moreous system (CMS) and myocardial), glomerulomephritis, especial acid molecule are infection, inflammation, inflammatic of a novel enzymatic cinfection. This sequence represents the substrate of a novel enzymatic
                                                                                                                                                              Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.
                                                                                                                                                                                                                                                                                                                                  describes an enzymatic nucleic acid molecule (I) which down
                                                     Stinchcomb DT, Mcswiggen J, Draper KG,
                                                                                                                                                                                                                                                                               Claim 3; Page 34; 72pp; English.
                                                                                                           WPI; 2003-340953/32.
(DRAP/) DRAPER K G.
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Sequence 17 BP; 0 A; 6 C; 9 G; 0 T; 2 U; 0 Other;

Query Match
3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels 305 GAGCCCCGGGGACCGC 320 ò

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0; Gaps

17 chácccccccccc 2 셤

ACA06586; ACA06586/c

NFKB sub-unit modulating inozyme substrate #405.

Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; peroreatic cancer; peroreatic cancer; peroreatic cancer; peroreatic cancer; peroreatic cancer; head and neck cancer; ovarian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; decetaxel; cisplatin; methorrexate; cyclophosphanide; docetaxel; cisplatin; methorrexate; cyclophosphanide; docetaxel; inflammatory disease; asthma; diabetes; pheumatoid arthritia; restenosis; Crohn's disease; asthma; diabetes; pheumatoid arthritia; restenosis; Crohn's disease; obesity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection; se

Homo sapiens

ACA06586 standard; RNA; 17 BP. 03-JUN-2003 (first entry) RESULT 477 

Draper KG; 92US-00987132. 94US-00245466. 94US-00291932. 96US-00777916. Mcswiggen J, 23-MAY-2001; 2001US-00864785. STINCHCOMB D T. MCSWIGGEN J. DRAPER K G. WPI; 2003-340953/32. US2002177568-A1 Stinchcomb DT, 15-AUG-1994; 07-DEC-1992; 18-MAY-1994; 23-DEC-1996; 28-NOV-2002 (STIN/) (DRAP/)

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.

Claim 3; Page 33; 72pp; English.

Sequence 17 BP; 1 A; 7 C; 6 G; 0 T; 3 U; 0 Other;

Gaps ö Query Match
3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels

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ACA09010 standard, RNA; 17 BP 266 GCACCTGGAGCAGGGC 281 17 dchgcrdchdchdddc 2 RESULT 478 ACA09010 %%%%%%%% 셤 ઠ

NFKB sub-unit modulating amberzyme substrate #173. (first entry) 03-JUN-2003

ACA09010;

Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme;

G-cleaver, amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; brostate cancer; colorectal cancer; brain cancer; carcer; carcer; carcer; cancer; cancer; cancer; cancer; carcer; carcer; cancer; cancer; cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemocherapy; paclitaxel; docetaxel; cisplatin; methorrexate; cyclophosphamide; doxorubin; fluorouracil carboplatin; edatrexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatoid arthritis; restensis; Crohn's disease; obseity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection; ss. 

Homo sapiens.

US2002177568-A1.

28-NOV-2002.

23-MAY-2001; 2001US-00864785

92US-00987132, 94US-00245466, 94US-00291932, 96US-00777916, 18-MAY-1994; 15-AUG-1994; 23-DEC-1996; 07-DEC-1992;

STIN/) STINCHCOMB D T. MCSWIGGEN J.

DRAPER K G. DRAP/)

Mcswiggen J, Draper KG; Stinchcomb DT,

WPI; 2003-340953/32.

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.

Claim 3; Page 54; 72pp; English.

The invention describes an enzymatic nucleic acid molecule (1) which down regulates expression of a sequence encoding a subunit of nuclear factor requisation. The enzymeatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A.

(1) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially MG2-f. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, cervical, head and name cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, or emiticable or radiation therapy. The enzymatic and antisense nucleic acid molecules are also useful for treating inflammatory disease such as theumatoid arthritis, restencisis, asthma, Crohn's disease, diabetes, celection, gene therapy applications, isohaemia/reperfusion injury (central nervous system (CNS) and myocardial), glomerulonephritis, sequence represents the substrate of a novel enzymatic acid molecule. This sequence represents the substrate of a novel enzymatic

Sequence 17 BP; 4 A; 4 C; 8 G; 0 T; 1 U; 0 Other;

Gaps . Query Match 3.0%; Score 12.8; DB 1; Length 17; Best Local Similarity 81.2%; Pred. No. 3.2e+02; Matches 13; Conservative 1; Mismatches 2; Indels

286 CCAAGCTGGTGAAGGA 301 

2 CCAGGCUGGGGAAGGA 17

RESULT 479 ADA99410

ADA99410 standard; DNA; 17

ADA99410;

20-NOV-2003 (first entry)

Human MDZ3 scanning oligonucleotide SEQ ID 399.

Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MD23; MD21; MD212; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.

Homo sapiens.

EP1281758-A2.

30-JUL-2002; 2002EP-00016874.

02-AUG-2001; 2001US-00922181

(AEOM-) AEOMICA INC.

Shannon M, Gu Y, Nguyen C;

WPI; 2003-423107/40.

New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.

Example 8; SEQ ID NO 399; 103pp; English.

The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ12, MDZ3 is canceded at chromosome 7422.1, MDZ4 is encoded at chromosome 6921.3-22.2, MDZ7 is encoded at chromosome 602.2 is encoded at chromosome 602.3 in an expression of mDZ3, MDZ4, MDZ3, 

Sequence 17 BP; 3 A; 7 C; 1 G; 6 T; 0 U; 0 Other;

Gaps ö 3.0%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 3.2e+02; tive 0; Mismatches 2; Indels 14; Conservative Query Match Best Local Similarity Matches

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RESULT 480

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ABZ61658 standard; RNA; 17 BP. ABZ61658

ABZ61658;

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Page 233

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Macejak D,
Roberts B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-229207/22.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LEE P.
DRAPER K.
ROBERTS E.
                                                                                  Hepatitis C virus.
                                                                                                                                            WO200281494-A1
                                                                                                                                                                                                   17-0CT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Blatt L, M
Draper K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (RIBO-) | (BLAT/) | (MACE/) | (MCSW/) | (MCRK/) | (PAVC/) | (LEEP/) | (DRAP/) | (CRAP/) | (CRAP/) | (ROBE/) | (ROBE/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 482
ACD58724/c
ID ACD587
XX
AC ACD587
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Loca
Matches
d
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  δ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule act useful for reducing HER2, K-Ras, H-Ras, and HIV acid molecules are useful for treating breast, ovarian, colorectal, lung, prostate, also useful for treating breast, ovarian, colorectal, lung, prostate, shown in NaZS9889 - ABZ62216, ABZ64544 - ABZ65531, ABZ65200 - ABZ65214, ABZ65531, ABZ65310 - ABZ65524,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleic acid molecule, Hepatitis C virus, HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; hammerhead riboxyme; DNAzyme; inozyme; amberzyme; G-cleaver riboxyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
                                                                                                                                      Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras; enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV; anti-rheumatic; cancer; AIDS; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17 BP; 6 A; 3 C; 6 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HCV DNAzyme substrate sequence #946.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 58; Page 119; 185pp; English.
                                                                                              Human H-Ras DNAzyme target #449.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 481
ACD58640/c
ID ACD58640 standard; RNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      287 CAAGCTGGTGAAGGAC 302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-MAY-2001; 2001US-0294140P.
06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-MAY-2002; 2002WO-US016840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ribozymes of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-140484/13.
                                                                                                                                                                                                                                                                                                                           WO200297114-A2.
                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mcswiggen J;
                                       21-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                     05-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ACD58640;
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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HHV) or Honocleic acid molecules include antisense and enzymatic nucleic acids such as harmerhead ribozymes, DNAzymes, or inozymes, zinzymes, amberzymes, and G-cleaver ribozymes, Also disclosed are nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptase primer sequences, as well transcriptase and/or HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV CC as oligonucleotides that specifically bind the Enhancer I region of HBV Genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds methods of the invention are useful for the treatment of degenerative and disease states related to HBV and HV virfection, replication and gene certains are active the present sequence espression such as cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence espressent in the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ä
liver failure, hepatocellular carcinoma, hepatotropic, cytostatic; virucide, antiinflammatory, substrate; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Lee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
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87.5%; Pred. No. 3.2e+02;
ative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Morrissey D, Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 1 A; 8 C; 4 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Page 250; 387pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        77 GGGCCGCGCAGTGGAC 92
                                                                                                                                                                                                                                                                                                                                                                            26-MAR-2001; 2001US-00817879.
08-UIN-2001; 2001US-0087478.
08-UIN-2001; 2001US-0296876P.
24-OCT-2001; 2001US-0335059P.
05-DEC-2001; 2001US-0337055P.
                                                                                                                                                                                                                                                                                                                  26-MAR-2002; 2002WO-US009187.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RIBOZYME PHARM INC.
BLATT L.
MACEGJAK D.
MCSWIGGEN J.
MORRISSEY D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACD58724 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14; Conservative
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Nucleic acid molecule, Hepatitis C virus, HCV; Hepatitis B virus, HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid, hammerhead ribozyme; inozyme; inozyme; inozyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative, disease state, HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
                                                                                                                                                                                                                                                                                                                                              á
                                                                                                                                                                                                                                                                                                                                              Mcswiggen J, Morrissey D, Pavco P,
                          HCV DNAzyme substrate sequence #974.
                                                                                                                                                                                                          08-JUN-2001; 2001US-00877478.
08-JUN-2001; 2001US-0296876P.
28-CVT-2001; 2001US-0335059P.
05-DEC-2001; 2001US-0337055P.
                                                                                                                                                                                26-MAR-2002; 2002WO-US009187.
                                                                                                                                                                                                  26-MAR-2001; 2001US-00817879.
                                                                                                                                                                                                                                                     RIBOZYME PHARM INC.
BLAIT 1.
        24-SEP-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                             Macejak D,
Roberts E;
                                                                                                                                                                                                                                                                 BLATT I.
MACEJAK D.
MCSWIGGEN J.
MORRISSEY D.
PAUCO P.
                                                                                                                          Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                             ROBERTS E.
                                                                                                                                                                                                                                                                                                          LEE P.
DRAPER K.
                                                                                                                                            WO200281494-A1.
                                                                                                                                                                                                                                                                                                                                      Blatt L, Me
                                                                                                                                                              17-0CT-2002.
                                                                                                                                                                                                                                                      (RIBO-)
(BLAT/)
(MACE/)
(MCSW/)
(MORR/)
(PAVC/)
(LEEP/)
(DRAP/)
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Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus nfection.

WPI; 2003-229207/22.

Claim 1; Page 251; 387pp; English.

The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as harmerhead ribozymes, DNAzymes, and enzymatic nucleic acids such as harmerhead ribozymes, DNAzymes, or inozymes, zinzymes, amberzymes, and G-cleaver ribozymes, Also disclosed are nucleic acid decoy molecules and G-cleaver ribozymes, Also disclosed transcriptase and/or HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and methods of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene carcinoma. The present sequence represente a substrate for one of the HCV invertion.

C DNAzyme or minus strand DNAzyme sequences disclosed in the present

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Sequence 17 BP; 2 A; 7 C; 5 G; 0 T; 3 U; 0 Other;

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Gaps
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1.0%; Score 12.8; DB 1; Length 17; larity 87.5%; Pred. No. 3.2e+02; Conservative 0; Mismatches 2; Indels
Query Match
Best Local Similarity
Matches 14; Conserva
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The invention relates to a nucleic acid molecule which encodes a Na+/H+ exchanger like protein (NHELPI). The NHELPI nucleic acid molecule, NHELPI polypeptide, an antibody against the protein or its antigen-binding fragment is useful in therapy. The NHELPI nucleic acid molecule, NHELPI polypeptide and an agonist are particularly useful for manufacturing a medicament for treating or preventing a disorder associated with antigen-binding fragment, and an anteagonist, are useful for manufacturing a medicament for treating or preventing a disorder associated with increased expression or activity of human NHELPI. The NHELPI for manufacturing increased expression or activity of human NHELPI. The NHELPI molesic acid or protein is useful as passive replacement therapy, as a vaccine, or in diagnostic methods. This sequence corresponds to a 17-mer oligonucleotide spanning the sequence of the human NHELPI gene (ADC03514).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New human sodium-hydrogen exchanger like protein 1 (NHELP1), useful as passive replacement therapy or as a vaccine for treating or preventing disorders associated with aberrant expression or activity of human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                          ss; gene therapy; vaccine; sodium/hydrogen exchanger like protein; NHELP1; passive replacement therapy; vaccine; diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                     Human Na/H exchanger-like protein 1 gene oligonucleotide #702
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 3.0%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 3.2e+02; Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 2 A; 5 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 2; SEQ ID NO 742; 468pp; English.
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ADC04256/c
ID ADC04256 standard; DNA; 17 BP.
XX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-JAN-2001; 2001WO-USG00666.
23-MAY-2001; 2001US-00864761.
21-DEC-2001; 2001US-0343331P.
                                                                                                                            BP.
310 CCGGGGACCGCGTGCT 325
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           39 GAAGATGGCCACCACT 54
                                                                                                                                                                                                                                                                                                                                                                                                                                                25-JAN-2002; 2002EP-00001160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GAAAATGGCCAGCACT 2
                                 16 ccccccccccicaticar 1
                                                                                                                          ADC04255 standard; DNA; 17
                                                                                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (AEOM-) AEOMICA INC
                                                                                                                                                                                                                                                                                                                                     Homo sapiens,
                                                                                                                                                                                                                                                                                                                                                                       EP1273660-A2
                                                                                                                                                                                                     18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                          08-JAN-2003.
                                                                                                                                                                 ADC04255;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NHELP1.
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The invention relates to a nucleic acid molecule which encodes a Na+/H+ exchanger like protein (NHELP1). The NHELP1 nucleic acid molecule, NHELP1 polypeptide, an antibody against the protein of its antibading fragment is useful in therapy. The NHELP1 nucleic acid molecule, NHELP1 polypeptide and an agonist are particularly useful for manufacturing a medicament for treating or preventing a disorder associated with decreased expression or activity of human NHELP1. The antibody or its antigen-binding fragment, and an antagonist, are useful for manufacturing a medicament for treating or preventing a disorder associated with increased expression or activity of human NHELP1. The NHELP1 andacturing a nuclean structuring a medicament for treating or preventing a disorder associated with or protein is useful as passive replacement therapy, as a vaccine, or in diagnostic methods. This sequence corresponds to a 1-mer oligonucleotide spanning the sequence of the human NHELP1 gene (ADC03514).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   plant growth; plant growth trait modulation; Brassicaceae; Arabidopsis; Brassica; Zea; Oryza; Triticum; Hordeum; Lolium; Sorghum; Glycine; Medicago; Helianthus; Lactuca; Beta; Vitis; Solanum; Lycopersicon; Capsicum; Gossypium; Hevea; Linum; Prunus; Citrus; Populus; Pinus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New human sodium-hydrogen exchanger like protein 1 (NHELP1), useful as passive replacement therapy or as a vaccine for treating or preventing disorders associated with aberrant expression or activity of human
                                                                                                                 ss; gene therapy; vaccine; sodium/hydrogen exchanger like protein; NHELP1; passive replacement therapy; vaccine; diagnosis.
                                                                            Human Na/H exchanger-like protein 1 gene oligonucleotide #703.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ouery Match 3.0%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 3.2e+02; Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Plant growth associated polynucleotide seg id 203.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 2 A; S C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 2; SEQ ID NO 743; 468pp; English.
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                                                                                                                                                                                                                                                                                                                                           30-JAN-2001; 2001WO-US000666.
23-MAY-2001; 2001US-00864761.
21-DEC-2001; 2001US-0343331P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16 GAAAATGGCCAGCACT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-JAN-2004 (first entry)
                                       18-DEC-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                               (AEOM-) AEOMICA INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-302724/30.
                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                       EP1273660-A2.
                                                                                                                                                                                                                                                                 08-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADE25228;
  ADC04256;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gu Y;
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MAKA KEKAKAKA
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                                                                                                                                                                                                                                                                                                                                         The invention describes an isolated or recombinant polypeptide (I) comprising a sequence: (a) comprising 1 of 30 sequences (S1), as given in the specification; or a conservative variant; (b) encoded by 1 of 30 sequences (S2), as given in the specification, or a conservative variant; (c) encoded by a sequence that hybridises under stringent conditions to activity of (I) is modulated to modulate a plant growth trait in a city Arabidopsis, Brassica, Zea, Orac, Triticum, Hordeum, Lolium, Sorghum, Glycine, Medicago, Helianthus, Lactuca, Beta, Vitis, Solanum, Lycopersicon, Capsicum, Gossypium, Hevea, Linum, Prunus, Citrus, Populus, Pycopersicon, Capsicum, Gossypium, Hevea, Linum, Prunus, Citrus, Populus, growth trait. This sequence represents a polynucleotide isolated from the plant growth associated genes of the invention that can be used a a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Probe, component B, promoter; human; signal peptide; primer; RACE; low molecular weight protein; urine; TGF-alpha; receptor; amplify; inflammation; coagulation; tumour; anglogenesis; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Match 3.0%; Score 12.8; DB 1; Length 17; Local Similarity 87.5%; Pred. No. 3.2e+02; les 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17 BP; 6 A; 2 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                      Example 2; SEQ ID NO 203; 81pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       primer, probe or genetic marker.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Component B gene primer, CKCB2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ87873 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        285 ACCAAGCTGGTGAAGG 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          93WO-EP003645.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2 ATCAAACTGGTGAAGG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21-DEC-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAR-2003
27-JUL-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9414959-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-JUL-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAQ87873;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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AAQ87873/c
                                                                                                                                                                                                                                                                                            Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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New isolated or recombinant polypeptide for use in modulating a plant growth trait in a flowering plant e.g. in Arabidopsis, Brassica, Zea, or

Bowen BA, Haudenschild CD, Buckler ES;

WPI; 2003-803305/75.

(LYNX-) LYNX THERAPEUTICS INC.

07-JAN-2003; 2003US-00338777. 09-JAN-2002; 2002US-0347288P.

US2003188343-A1 Magnoliophyta.

02-OCT-2003

The sequences given in AAQB7870-75 are primers which were used in the amplification of the component B CDNA. These primers were used in the carpid annolation of the component B CDNA. These primers were used in the carpid annolation of the gene including exon 2 and the poly-A tail. The component B cegons of the gene including exon 2 and the poly-A tail. The component B cegons of the gene contains three exons and two introns. Exon 1 is 84 bp and contains cegons of the responsible of the material contains of the material contains of the material of the ma New protein, component B, isolated from urine - with antiinflammatory, anticoagulant and anti-tumour activities, also related nucleic acid, vectors and transformed cells. (ISTF ) ARS APPLIED RES SYST HOLDING NV. Example 4; Page 28; 55pp; English. WPI; 1994-234696/28. 22-DEC-1992; AAA58496; Sirna A; RESULT 487 AAA58496 쉱 ò

88. Gaps BLM gene cluster; bleomycin gene cluster; polyketide metabolite; bleomycin; bleomycin analogue; holo-carrier protein; thiazolidine; thiazoline; bithiazoline; microbial metabolite; sugar; PCR primer; , 0 PCR primer used to amplify bleomycin (BLM) gene cluster ORF19. Query Match 3.0%; Score 12.8; DB 1; Length 18; Best Local Similarity 87.5%; Pred. No. 3.6e+02; Matches 14; Conservative 0; Mismatches 2; Indels Sequence 18 BP; 3 A; 7 C; 5 G; 3 T; 0 U; 0 Other; 285 ACCAAGCTGGTGAAGG 300 AAAS8496 standard; DNA; 18 BP 06-JAN-1999; 99US-0115435P. 05-FEB-1999; 99US-0118848P. 05-JAN-2000; 2000US-00477962. 06-JAN-2000; 2000WO-US000445 17 ACCACGCTGGTGACGG 2 20-OCT-2000 (first entry) Streptomyces verticillus WO200040704-A1. 13-JUL-2000. 

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PCR primers AAAS8474-A58541 were used to amplify open reading frames (ORFs) 8 to 41 of the BLM (Bleomycin) gene cluster. The proteins encoded by the gene cluster are useful for producing peptides and/or polyketide metabolites, especially bleomycin or bleomycin analogues. They are also useful for chemically modifying biological molecules to produce branched methyl groups, and for coupling amino acids and fatty acids. They may be reacted with an apo-carrier protein and coenzyme A to produce a holocarrier protein. The BLM gene cluster or catalytic domains can be used hidividually or collectively to produce thiazolidine, thiazoline, bithiazoline and bithiazoline—containing microbial metabolites. The BLM gene cluster may also be used to produce sugars
                                                                                                                       New bleomycin gene cluster components useful for peptide and/or polyketide metabolites, especially bleomycin, production and for chemically modifying biological molecules.
                                      Edwards DJ;
                                                                                                                                                                                                                  Disclosure; Page 22; 162pp; English
                                        Chen M,
                                      Du L, Sanchez C,
                                                                                   WPI; 2000-465974/40.
                                        Shen B,
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(REGC ) UNIV CALIFORNIA

92IT-RM000919

294 GTGAAGGACCTGAGCC 309 1 GTGAAGGACCTCGGCC 16 셤 ठ

Query Match
Best Local Similarity 87.5%; Pred. No. 3.6e+02;
Matches 14; Conservative 0; Mismatches 2; Indels

Sequence 18 BP; 3 A; 5 C; 8 G; 2 T; 0 U; 0 Other;

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0; Gaps

AAH40454 standard; DNA; 18 14-AUG-2001 (first entry) AAH40454; RESULT 488 AAH40454/c

SNP specific lower PCR primer SEQ ID 3250.

Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPB; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; ss.

15-OCT-1999; 99US-0160096P. 13-OCT-2000; 2000WO-US028436. WO200129262-A2 Homo sapiens. 26-APR-2001. 

(ORCH-) ORCHID BIOSCIENCES INC. Piccult-Newburg L, Pohl M; WPI; 2001-290930/30.

New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polymucleotide polymorphism in a nucleic acid sample.

Claim 1; Page 66; 83pp; English

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Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide polymorphisms SNPs. The present invention cities of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the includes kits for determining the presence or absence of a SNP, using the cligonucleotides of the invention. The PCR primers are used to amplify a SC pligonucleotides of the invention. The PCR primers are used to amplify a cligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The performing a single-nucleotide primer extension reaction. The cligonucleotides are useful for determining the presence, absence or dentity of a SNP and for genotyping nucleic acid samples, for e.g. to assess by association analysis the genotype of an individual or group of individuals, having a pathological phenotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. or agammaglobulinemia, diabetes insipidus, lesch Nyhan syndrome, muscular dystrophy, familial hypercholesterolaemia polycystic kidney disease, osteogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial clisease of which a component is or may be genetic such as autoimmune diseases, including, rheumatoid arthritis, multiple sclerosis, inflammation, cancer, nervous system diseases and infection and containing DNA sequence represents a PCR primer specific containing DNA sequence

Sequence 18 BP; 3 A; 8 C; 5 G; 2 T; 0 U; 0 Other;

0; Gaps Match 3.0%; Score 12.8; DB 1; Length 18; Local Similarity 87.5%; Pred. No. 3.6e+02; les 14; Conservative 0; Mismatches 2; Indels 299 GGACCTGAGCCCCGGG 314 Query Match Best Loca Matches ઠે

**é**GTCCTGAGCCCAGGG 18

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ABL40174 standard; DNA; 18 BP RESULT 489 ABL40174 

(first entry) 21-MAY-2002 ABL40174;

Mouse reelin protein CR-50 epitope region PCR primer SEQ ID NO:11.

Mouse; realin protein CR-50 epitope region; elucidation; neuron; cerebral disturbance; reelin protein; neuroprotective; PCR primer; ss.

Mus musculus.

JP2002017361-A.

22-JAN-2002.

04-JUL-2000; 2000JP-00202801.

04-JUL-2000; 2000JP-00202801.

(RIKE ) RIKEN KK.

WPI; 2002-221707/28.

Reelin protein CR-50 epitope region, useful for diagnosis and treatment of cerebral disturbance.

Example 2; Page 7; 16pp; Japanese.

The present invention describes the mouse realin protein CR-50 epitope region, which contains the CR-50 antibody recognition site and is free from Fepordin domains and repetitive sites. Also described are: (1) an expression vector comprising a polymucleotide encoding a realin protein epitope region; (2) host cells with transfected the expression vector;

(3) polypeptides prepared by culture of the host cells; and (4) polynucleotides comprising the 351 base sequence given in ABE40165 which pencedes the 117 amino acid sequence given in ABE06244; and (5) use of the polynucleotide for diagnosis and/or treatment of diseases caused by abnormal positioning of neural cells, and stimulation of association of reelin protein. The mouse reelin protein CR-50 epitope region has neuroprotective activity, and can be used in the diagnosis and treatment of cerebral disturbance due to an abnormal reelin gene and positioning of neurons. The present sequence represents a PCR primer for the mouse reelin protein CR-50 epitope region, which is used in an example from the present invention 888888888888888

Sequence 18 BP; 4 A; 5 C; 6 G; 3 T; 0 U; 0 Other;

Gaps 3.0%; Score 12.8; DB 1; Length 18; 87.5%; Pred. No. 3.6e+02; Azive 0; Mismatches 2; Indels 0; Best Local Similarity 87.5 Matches 14; Conservative Query Match

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7438/c ABK27438 standard; DNA; 18

ABK27438;

09-APR-2002 (first entry)

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Colon cancer associated cDNA CATX-7, 5' PCR primer.

Human; colon cancer; tumour; abnormal cell growth; melanoma; cervical cancer; colorectal adenocarcinoma; Wilms' tumour; leukaemia; lymphoma; antisense therapy; CATX; probe; primer; ss.

Homo sapiens.

WO200111047-A2.

15-FEB-2001.

08-AUG-2000; 2000WO-US021606.

09-AUG-1999; 99US-0147933P.

(FARE ) BAYER CORP.

Bowman BM, Wang K;

WPI; 2002-121548/16.

New isolated nucleic acid involved in growth regulation in human colonic epithelial cells, termed CATX, for diagnosing and treating abnormal cell growth, and for use as a probe/primer for detecting tumors.

Example, Page 88; 130pp, English.

The invention relates to an isolated nucleic acid (1) involved in growth regulation in human colonic epithelial cells, termed CATX. (1) is useful as a probe-primer for detecting tumours, preferably colon cancer. The nucleic acid, encoded polypeptide and antibody are useful in diagnosis and treatment of abnormal cell growth (such as cervical cancer, mealsonmas, colorectal adenocarcinomas, wilms' tumour, leukaemias and lymphomas), in screening assays for the treatment of abnormal cell growth, for raising antibodies, and to screen for peptide analogues and antagonists. (1) is useful as a biomarker for human tumour cells, e.g., colon cancer cells, for generating probes and primers designed for identifying and/or cloning homologues in other cell types, in antisense largament, so that premaismant cells can be identified prior to their spreading throughout the human body. (I) allows early detection 

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of potentially cancerous conditions, and treatment of the cancerous conditions prior to spread of the cancer cells throughout the body, or prior to development of an irreversible cancerous condition. ABXC7426-ABX27469 represent human colon cancer associated coding sequences and
                                                                                                       primers of the invention
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Sequence 18 BP; 3 A; 5 C; 9 G; 1 T; 0 U; 0 Other;

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Query Match

3.0%; Score 12.8; DB 1; Length 18;
Best Local Similarity 87.5%; Pred. No. 3.6e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                338 CCAGGGCCGGCTGCTC 353
                                                                                                          18 CCAGGGCTGGCTCCTC 3
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ABK27436 standard; DNA; 18 BP 09-APR-2002 (first entry) ABK27436; 

Colon cancer associated cDNA CATX-6, 5' PCR primer.

Human; colon cancer; tumour; abnormal cell growth; melanoma; cervical cancer; colorectal adenocarcinoma; Wilms' tumour; leukaemia; lymphoma; antisense therapy; CATX; probe; primer; ss.

Homo sapiens,

WO200111047-A2.

15-FEB-2001.

08-AUG-2000; 2000WO-US021606

99US-0147933P 09-AUG-1999;

(FARB ) BAYER CORP.

Bowman BM, Wang K;

WPI; 2002-121548/16.

New isolated nucleic acid involved in growth regulation in human colonic epithelial cells, termed CATX, for diagnosing and treating abnormal cell growth, and for use as a probe/primer for detecting tumors.

Example; Page 88; 130pp; English.

The invention relates to an isolated nucleic acid (I) involved in growth regulation in human colonic epithelial cells, termed CATX. (I) is useful as a probe/primer for detecting tumours, preferably colon cancer. The nucleic acid, encoded polypeptide and antibody are useful in diagnosis and treatment of abnormal cell growth (such as cervical cancer, lewaemias and Lymphomas), in screening assays for the treatment of abnormal cell growth, for raising antibodies, and to screen for peptide analogues and antagonists. (I) is useful as a biomarker for human tumour cells, e.g., colon cancer cells, for generating probes and primers designed for identifying and/or cloning homologues in other cell types, in antisense therapy, and in tissue profiling. (I) identifies cancer cells at an early stage of development, so that premalignant cells and piners cells at an early cape of development, so that premalignant cells and incompleted prior to their spreading throughout the human body. (I) allows early detection of potentially cancerous conditions prior to spread of the cancer cells throughout the body, or prior to development of an irreversible cancerous condition. ABKZ7456-ABKZ7459 represent human colon cancer associated coding sequences and primers of the invention

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Query Match
3.0%; Score 12.8; DB 1; Length 18;
Best Local Similarity 87.5%; Pred. No. 3.6e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
                                                                               338 CCAGGGCCGGCTGCTC 353
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Seguence 18 BP; 3 A; 5 C; 9 G; 1 T; 0 U; 0 Other;

'432/c ABK27432 standard; DNA; 18 ABK27432; RESULT 492 ABK27432/c

BP.

09-APR-2002 (first entry)

Colon cancer associated cDNA CATX-4, 5' PCR primer.

Human, colon cancer, tumour, abnormal cell growth, melanoma; cervical cancer, colorectal adenocarcinoma; Wilms' tumour; leukaemia, lymphoma; antisense therapy; CATX; probe; primer; 88.

Homo sapiens

WO200111047-A2.

15-FEB-2001.

08-AUG-2000; 2000WO-US021606.

09-AUG-1999; 99US-0147933P

(FARB ) BAYER CORP.

Bowman BM, Wang K,

WPI; 2002-121548/16.

New isolated nucleic acid involved in growth regulation in human colonic epithelial cells, termed CATX, for diagnosing and treating abnormal cell growth, and for use as a probe/primer for detecting tumors.

Example, Page 87; 130pp; English.

The invention relates to an isolated nucleic acid (I) involved in growth regulation in human colonic epithelial cells, termed CATX. (I) is useful as a probe/primer for detecting tumours, preferably colon cancer. The nucleic acid, encoded polypeptide and antibody are useful in diagnosis and treatment of abnormal cell growth stumour, leukaemias and and treatment of abnormal cell growth, to raising antibodies, and to screen for peptide analogues and growth, for raising antibodies, and to screen for peptide analogues and growth, for raising antibodies, and to screen for peptide analogues and contagonists. (I) is useful as a biomarker for human tumour cells, e.g., colon cancer cells, for generating probes and primers designed for identifying and/or cloning homologues in other cell types, in antisense therapy, and in tissue profiling. (I) identifies cancer cells at an early crappy and in tissue profiling. (I) identifies cancer cells at an early crappy and in to spread of the cancer cells and early conting spreading throughout the human body. (I) allows early detection of potentially cancerous conditions and treatment of the cancerous conditions prior to spread of the cancer cells throughout the human potential conditions prior to spread of the cancer cells throughout the hody. Or prior to development of an irreversible cancerous condition. ABK27426-CC primers of the invention 

Sequence 18 BP; 3 A; 5 C; 9 G; 1 T; 0 U; 0 Other;

Gaps ö Query Match 3.0%; Score 12.8; DB 1; Length 18; Best Local Similarity 87.5%; Pred. No. 3.6e+02; Matches 14; Conservative 0; Mismatches 2; Indels

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The present invention describes a protein which is fused with a monoclonal antibody against an antigen present on cell surface and which can transfer a gene by combining with the gene and containing a human type single-stranded monoclonal antibody and a peptide which is the combining site for the gene. Also described is a complex of a monoclonal antibody-fused protein which is a complex of monoclonal antibody-fused protein and a math a method for the preparation of a monoclonal antibody-fused protein against a receptor present on cell surface in antibody-fused protein against a receptor present on cell surface in such a minoclonal antibody against a receptor present on cell surface is used as the template to amplify a single-stranded antibody gene of a course type monoclonal antibody by PCR; (2) the framework portion of the mouse type monoclonal antibody by PCR; (2) the framework portion of the antibody gene of a human type monoclonal antibody is converted to prepare a single-stranded immunoporter gene; and (4) the human type single-stranded immunoporter gene; and in a microbe to prepare a recombinant protein of the human type single-stranded immunoporter. Also described is a method for introducing the above complex of monoclonal antibody-fused protein of the through a cell surface receptor. The method is used for the preparation of surface. The present sequence represents an oligonucleotide which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                         Monoclonal antibody; fusion protein; antigen; cell surface; receptor; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A protein fused with a monoclonal antibody against an antigen present on
                                                                                                                                                                                                                                                                                 Monoclonal antibody related oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example, Page 6; 24pp; Japanese.
                                                                                                                                                  ABA94181 standard; DNA; 18 BP.
338 CCAGGGCCGGCTGCTC 353
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29-MAY-2000; 2000JP-00158575.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cell surfaces
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                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
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Sequence 18 BP; 4 A; 5 C; 7 G; 2 T; 0 U; 0 Other;
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Gaps

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Query Match 3.0%; Score 12.8; DB 1; Length 18; Best Local Similarity 87.5%; Pred. No. 3.6e+02; Matches 14; Conservative 0; Mismatches 2; Indels

293 GGTGAAGGACCTGAGC 308

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Sequence 18 BP; 4 A; 5 C; 7 G; 2 T; 0 U; 0 Other;

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GGTGCACCTGGAGCAG 278
                   ścreckecreckecke 18
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263
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Query Match 3.0%; Score 12.8; DB 1; Length 18; Best Local Similarity 87.5%; Pred. No. 3.6e+02; Matches 14; Conservative 0; Mismatches 2; Indel8

Human; growth inhibitory gene; retinoid; retinoic acid response element; RARE site; therapy; promyelocytic leukaemia; cancer chemoprevention; cytostatic; secreted cell adhesion protein beta IG-H3 promoter; PCR primer; ss.

Homo sapiens

Human beta IG-H3 promoter DNA amplifying antisense PCR primer.

12-MAR-2002 (first entry)

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Gaps ö

AAD24955;

AAD24955 standard; DNA; 18 BP.

RESULT 494

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The present invention relates to human chondroitin 6-sulfotransferase (C6ST) proteins and polynucleotides encoding such proteins. Sequences of the invention are useful in the molecular study of human extracellular matrix, for studying the biological functions of chondroitin 6-sulphate (C6S), in screening test for detecting C6ST polymorphs, for ascertaining and evaluating the role C6ST plays in atherosclerosis and for identifying potential therapeutics, i.e., inhibitors of enzyme or modulators of gene expression. The present DNA sequence is a PCR primer which is used for amplifying human C6ST gene
                                                                                                                                        Human, chondroitin 6-sulfotransferase, C6ST, chondroitin 6-sulphate, C6S, biological function; extracellular matrix; atherosclerosis; therapeutic; gene expression; enzyme; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel recombinant human chondroitin 6-sulfotransferase polynucleotide segment, useful in molecular study of human extracellular matrix, and for studying biological functions of chondroitin 6-sulfate.
                                                                                                           Human C6ST gene amplifying 5' PCR primer #3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Col 17; 15pp; English.
                                                                                                                                                                                                                                                                                                                                                                                        (UYJE-) UNIV JEFFERSON THOMAS
               AAD41288 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                                         98US-00015188.
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                                                                                                                                                                                                                                           US6399358-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                       Williams KJ,
                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                         29-JAN-1998;
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02-JUL-1997;
                                                                          30-OCT-2002
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                                            AAD41288;
AAD41288
ID AAD4
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Tang

Cheng Y,

Shen B,

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The patent discloses growth inhibitory genes induced by retinoids. The invention also relates to recombinant expression constructs that express a reporter gene under the transcriptional control of a promoter for a gene which is expressed by retinoid induction. The promoter does not contain a retinoic acid response elements (RARE) site. The invention that her tealstes to reagents and methods for identifying compounds other than retinoids that modulate the expression of cellular genes. These compounds are useful for treating cancers such as promyelocytic leukaemia and cancer chemoprevention. The present DNA sequence is a PCR primer which is used for amplifying human secreted cell adhesion protein beta IG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Leinamycin biosynthesis gene cluster; Lmn; open reading frame; ORF; anti-tumour antibiotic; broad spectrum antimicrobial activity; dram-negative; fram-negative bacteria; chemical modification; metabolite; apo-carrier protein; holo-carrier protein; tumour; polyketide; hybrid polypeptide/polyketide metabolite; Inm production; cytostatic;
                                                                                                                                                                                                                                                          Expression construct encoding cellular genes, under control of a promoter regulated by retinoids and cells comprising the construct for identifying compounds that induce expression of the genes useful in treating cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Match 3.0%; Score 12.8; DB 1; Length 18; Local Similarity 87.5%; Pred. No. 3.6e+02; Loservative 0; Mismatches 2; Indels 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR primer #1 for S. atroolivaceus leinamycin gene cluster ORF lnmM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 18 BP, 4 A, 4 C, 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                          ñ
                                                                                                                                                                                            Chang
                                                                                                                                                                                                                                                                                                                                 Example 3; Page 27; 64pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       366 CTCACTTTCCTGGACC 381
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(KYOW ) KYOWA HAKKO KOGYO KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-MAR-2002; 2002WO-US008937
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                                                                                  25-MAY-2001; 2001WO-US017161.
                                                                                                                     26-MAY-2000; 2000US-0207535P.
                                                                                                                                                                                         Dokmanovic M,
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                                                                                                                                                     (UNII ) UNIV ILLINOIS FOUND
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                                                                                                                                                                                                                            WPI; 2002-075474/10
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              WO200192578-A2
                                                                                                                                                                                          Roninson IB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3-OCT-2002.
                                                  06-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABX34384;
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The present invention relates to the isolation of the Streptomyces atroolivaceus leinamycin (Lnm) biosynthesis gene cluster containing 71 cpn reading frames (ORFS) (ORFS) distrough -1.0 ORFS inma through lnm2, and ORFS +1 through +9). Leinamycin is a novel anti-tumour antibiotic containing an extinity against Gram-positive and Gram-mogative bacterum containing a method of produced by several Streptomyces species. It exhibits broad spectrum containing a method of the boat cell is a bacterium or enkaryotic cell, including a mammalian, yeast, plant, fungal, or insect cell. The molecule is an endogenous metabolite produced by the host cell. The molecule is an endogenous metabolite produced by the host cell. The molecule is an endogenous metabolite produced by the host cell. The molecule is an endogenous metabolite produced by the host cell. The sexpenously supplied containing a mino transferame. The polypeptides encoded by the Lnm gene cluster for converting an apo-carrier protein to a holo-carrier or are useful for converting an apo-carrier protein to a holo-carrier or making various peptide and/or polyketide, and/or hybrid compiled and/or polyketide and/or polyketide and/or hybrid constitution with other active domains to modify various target substrates. The Lnm gene cluster is useful to make various modified Lnm. Lnm, its analogue, or other polyketide, cendogenous man production to the proteins encoded by the ORFS are condogenous man production with other active domains to morphy condens and control polyketide and condens or their polyketide, or other polyketide, condens to make various modified Lnm. Lnm, its analogue, or other polyketide, condens to append the supplied and condens and production in cells and condens to make various modified Lnm. Lnm, its analogue, or other polyketide, condens an anumber of disorders depending upon the condogenous mand production and number of disorders dependent an anumber of disorders dependent an anumber of supply dependent and the supplication with the supplication with the
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                                                                                                               Novel gene cluster responsible for synthesis of leinamycin in
Streptomyces atroolivaceus useful for making various peptide and/or
polyketide, and/or hybrid polypeptide/polyketide metabolites.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3.0%; Score 12.8; DB 1; Length 18; 87.5%; Pred. No. 3.6e+02; rative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 18 BP; 3 A; 7 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                      Claim 1; Page 28; 185pp; English.
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Best Local Similarity 87.5'
Matches 14; Conservative
                                                                    WPI; 2003-018907/01.
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Med Apr 21 12:38:21 2004
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The present invention relates to the isolation of the Streptomyces atroolivaceus leinamycin (Lnm) biosynthesis gene cluster containing 71 open reading frames (ORFs) (ORFs) (ORFs - 15 through - 1, ORFs nm4 through lnm2, and ORFs + 1 through + 9). Leinamycin is a novel anti-tumour antibiotic produced by several Streptomyces species. It exhibits broad spectrum can information activity against Gram-positive and Gram-negative bacteria, but not against fungi. The polypeptides encoded by the Lnm biosynthesis content of the best cell is a bacterium or eukaryotic cell, including a mammalian, yeast, plant, fungal, or insect cell. The molecule in a mammalian, yeast, plant, fungal, or insect cell. The molecule is an endogenous metabolite produced by the host cell or exogenously supplied or an amino acid, and the polypeptide is a peptide synthetase or amino transferase. The polypeptides encoded by the Lnm gene cluster modules and/or catalytic domains are useful for converting an apo-carrier protein to a holo-carrier protein. Dum shows potent antitumour activity in tumnur models in vivo. The Lnm gene cluster modules and/or catalytic domains are useful for making various peptide metabolites. The proteins encoded by the ORFs are useful alone, or in combination with other active domains to modify various target substrates. The Lmm gene cluster is useful to upregulate endogenous Lnm production in cells and/or to make various modified Lnm. Lnm, its analogue, or other polyketide, peptide metabolites are useful as endogents. Created to permit Lnm production in cells and/or to treat a number of disorders, depending upon the type of metabolites. Abx3420-AAX4431 represent PCR primers used to amplify and individual ORFs of the S atroolivaceus leinamycin biosynthesis
                                                                                                                                                                                                                                                                                Novel gene cluster responsible for synthesis of leinamycin in
Streptomyces atroolivaceus useful for making various peptide and/or
polyketide, and/or hybrid polypeptide/polyketide metabolites.
                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; Page 29; 185pp; English.
22-MAR-2002; 2002WO-US008937.
                                                                                                  (REGC ) UNIV CALIFORNIA.
(KYOW ) KYOWA HAKKO KOGYO KK.
                                                   26-MAR-2001; 2001US-0278935P.
                                                                                                                                                                           Tang G;
                                                                                                                                                                                                                               WPI; 2003-018907/01.
                                                                                                                                                                              Cheng Y,
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                                                                                                                                                                              Shen B,
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Query Match
3.0%; Score 12.8; DB 1; Length 18;
Best Local Similarity 87.5%; Pred. No. 3.6e+02;
Matches 14; Conservative 0; Mismatches 2; Indels Seguence 18 BP; 4 A; 5 C; 7 G; 2 T; 0 U; 0 Other;

278 GGGCGCCACCAAGCTG 293 GAGCGCCCAAGCTG 18 ઠે

ABZ68636 standard; DNA; 18 BP. ABZ68636;

16-MAY-2003 (first entry) 8X8X8X8X8X8X8X8X8

Primer for extension of K121 antibody heavy chain variable region.

KI21 antibody; KI21-like antibody; kappa-type myeloma cell; kappa-type multiple myeloma; haematopoietic cell transplantation; apoptosis; kappa myeloma antigen; PCR; primer; ss.

Mus musculus.

WO2003004056-A1

16-JAN-2003 

06-JUL-2001; 2001AU-0006179

05-JUL-2002; 2002WO-AU000896

(PACM-) PACMAB PTY

Dunn RD, Raison RL,

WPI; 2003-210317/20.

Treating kappa-type multiple myeloma in a subject by administering a K121 -like antibody not conjugated to a toxin or a cytolytic agent. Example 8; Fig 9d; 65pp; English.

antibody heavy chain variable region. The primers were used to construct antibody heavy chain variable region. The primers were used to construct at X121-like antibody by oligonuclectide assembly using PCR. The K121-like antibody competes with K121 for binding to kappa-type myeloma cells. The K121-like antibody is used in the method of the invention. The specification describes a method for treating kappa-type multiple myeloma in a subject, comprising administering a K121-like antibody which is not conjugated to a toxin or a cytolytic agent. The method is useful for treating kappa-type multiple myeloma, autologous haematopoietic cell transplantation, killing kappa-type myeloma cells in a mixed population of cells and inducing apoptosis in kappa myeloma antigen (KMA) bearing

Seguence 18 BP; 4 A; 5 C; 7 G; 2 T; 0 U; 0 Other;

Gaps ·. 3.0%; Score 12.8; DB 1; Length 18; llarity 87.5%; Pred. No. 3.6e+02; Conservative 0; Mismatches 2; Indels Local Similarity es 14; Conservat Query Match Best Loca Matches

263 GGTGCACCTGGAGCAG 278 derechecheche 18

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ADD24785 standard; DNA; 18 BP. ADD24785; RESULT 499 ADD24785 

(first entry) 15-JAN-2004

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Gaps

; 0

Human CYP2D6 mutants G1661C and 1707delT probe H212.

diagnostic; pharmaceutical tolerance; side effect; drug; human; allelic variability; polymorphism; phase I; phase II; decoxification mechanism; PCR; primer; probe; NNT2; CYP2D6; CYP1A2; CYP3M4; mEH; TPNT; MTHFR; paraoxonase; CYP2C9; CYP2C19; CYP2E1; DPD; ss.

Homo sapiens

WO2003018837-A2.

06-MAR-2003.

22-AUG-2002; 2002WO-EP009386. 24-AUG-2001; 2001DE-01040651. 30-APR-2002; 2002DE-01019373.

(ADNA-) ADNAGEN AG.

Lustig M; Waschuetza S, Schnakenberg E,

WPI; 2003-290079/28

rage 417

Diagnostic kit, useful for assessing a subject's tolerance of drugs, comprises reagents for determining alleles of genes encoding detoxification enzymes.

Claim 6; Page 17; 156pp; German

This invention describes a novel diagnostic kit for determining tolerance of pharmaceuticals in humans by determining allelic variability of at least two polymorphisms of a human enzyme involved in phase I and/or I constitute two polymorphisms of a human enzyme involved in phase I and/or I constructed the detoxification mechanism in a blood, tissue or other human sample, where tolerance is determined from presence or absence of alleles. The kit comprises two pairs of agene for a human detoxification mechanism amplifies, by PCR, part of a gene for a human detoxification mechanism associated enzyme. The kit may also contain two further pairs of oligonucleotides, serving as probes for detection of amplified DNA segments, especially where the probes are complementary to a single strand of one allele of the target gene. The probes are labelled with segments, especially where the probes are complementary to a single strand of one allele of the target gene. The probes are labelled with construction. The enzymes detected include NAT2, CYP2D6, CYP3A4, mEH, TPMT, WHYER, paraxonaee, CYP2C9, CYP2C19, CYP2E1 or DPD.

CYP3A4, mEH, TPMT, WHYER, paraxonaee, CYP2C9, CYP2C19, CYP2E1 or DPD.

CYP3A4, mEH, TPMT a suitable dose and/or to predict if a subject will show xide-effects to a drug. The kit provides minimally invasive, safe and reliable determination of the metabolic capacity of phase I and/or II cenzymes at the molecular level. This sequence represents a probe used in constant of the kit of the invention.

Sequence 18 BP; 4 A; 3 C; 9 G; 2 T; 0 U; 0 Other;

Gaps ö Query Match 3.0%; Score 12.8; DB 1; Length 18; Best Local Similarity 87.5%; Pred. No. 3.6e+02; Matches 14; Conservative 0; Mismatches 2; Indels

15 CTGCGGGTGACCGAGG 30

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CAGTGGGTGACCGAGG 17

RESULT 500 ADE15233/c ID ADE15233 standard; DNA; 18 BP.

ADE15233; 

(first entry) 29-JAN-2004 Beer spoilage-associated primer SEQ ID 428.

ss; primer; detection; beer-spoilage; lactic acid bacteria; Gram-negative bacteria; spoilage bacteria.

Megasphaera cerevisiae

WO2002103043-A2.

27-DEC-2002.

19-JUN-2002; 2002WO-EP006808

19-JUN-2001; 2001DE-01029410.

(VERM-) VERMICON AG.

Snaidr J;

Beimfohr C,

WPI; 2003-175243/17.

New oligonucleotides, useful for rapid detection of beer-spoilage bacteria by in situ hybridization, are specific for type, genus or species

Claim 1; SEQ ID NO 428; 88pp; German.

This invention describes novel oligonucleotides used in a method for detecting beer-spoilage bacteria in a sample. The bacteria detected include lactic acid bacteria of the genera Lactobacillus or Pediococcus, especially the species L. coryniformis, L. perolens, L. buchneri, L. plantarum, L. fructivorans, L. lindneri, L. casei, L. brevis or P. damnosus or Gram-negative bacteria of the genera Pectinaus and Generalister, specifically P. frisingensis, P. cerevisiphilus and M. cerevisiae. The oligonucleotides of the invention provide rapid detection of spoilage bacteria (typically within 48 hours, compared with 7-12 days for conventional culture methods), can detect all relevant bacteria in parallel, can differentiate between species of the same genus, and are method of the invention.

Sequence 18 BP; 3 A; 7 C; 3 G; 5 T; 0 U; 0 Other;

Gaps ö / Match 3.0%; Score 12.8; DB 1; Length 18; Local Similarity 87.5%; Pred. No. 3.66+02; nes 14; Conservative 0; Mismatches 2; Indels Query Match

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RESULT 501 AAA27228 

AAA27228 standard; DNA; 19

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20-SEP-2000 (first entry)

AAA27228;

Forward PCR primer for FGF8.

Parkinson's disease; neurodegenerative disorder; PCR primer; FGF8; fibroblast growth factor 8; 8s.

Rattus sp.

WO200029550-A2.

25-MAY-2000.

99WO-US027613. 18-NOV-1999; 98US-00195569. 99US-00425462. 18-NOV-1998; 22-OCT-1999; (CALY ) CALIFORNIA INST OF TECHNOLOGY.

Studer L; Ceste M, Doyle J, Wold BJ, Mckay R,

WPI; 2000-387772/33.

Low oxygen culturing of central nervous system progenitor cells useful in treatment of neurodegenerative disorders.

Example 1; Page 36; 80pp; English.

A method for increasing the differentiation of undifferentiated central nervous system (CNS) cells in culture. This novel method involves culturing the cells in low ambient oxygen conditions. Differentiated cells can be used to treat neurodegenerative diseases such as Parkinson's disease. In order to determine the differentiated phenotype messenger RNA lusing PCR primers specific to certain genes. The present sequence is the forward PCR primer used to monitor the message level of FGF8

Sequence 19 BP; 5 A; 5 C; 7 G; 2 T; 0 U; 0 Other;

Mouse retinoid X receptor-gamma gene; RXR-gamma; exon E5; DNA binding domain; murine; transgenic animal; RXR-gamma knockout mouse; drug screening; reverse transcription-PCR; RT-PCR primer; 88.

(INRM ) INST NAT SANTE & RECH MEDICALE. (CNRS ) CENT NAT RECH SCI. (UYPA-) UNIV PASTEUR LOUIS. (BRIM ) BRISTOL-MYERS SQUIBB CO.

Chambon P, Kastner P; WPI; 2000-531490/48.

97US-00914256. 96US-0024175P.

19-AUG-1997; 19-AUG-1996;

25-JUL-2000. US6093873-A.

Mus sp.

Mouse retinoid X receptor-gamma gene exon B5 RT-PCR primer.

06-DEC-2000

AAA72197;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence is a PCR primer for the fibroblast growth factor 8 gene (FGP8). It was used in reverse transcription PCR to determine expression patterns of the FGF8 gene in cultured cells. These cells had been grown in low oxygen conditions, and had differentiated to form various types of neuronal cell. The different expression patterns can be used to determine which set of conditions promotes the differentiation of each type of neurone. The different cell types can be used for tissue transplantation, to treat disorders such as stroke, brain and spinal cord neurodegenerative disorders, Huntington's disease, other neurological disorders and psychiatric disorders parkinson's disease, neurological
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                                                                                                                                                                                                                                                                                                                      Rat; cell differentiation; neurodegenerative disorder; stroke; brain injury; spinal cord injury; Alzheimer's disease; epilepsy; Huntington's disease; Parkinson's disease; neurological disorder; cell transplantation; FGF8; fibroblast growth factor 8; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Culturing of neural crest stem cells useful for treatment of neurodegenerative disorders comprises culturing in low ambient oxygen conditions.
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 DB 1; Length 19;
                                  2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ceste M, Doyle J, Wold BJ, Morrison SJ, Anderson
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                                                                                                                                                                                                                                                                                     Fibroblast growth factor 8 mRNA PCR primer #1.
 3.0%; Score 12.8; DB 1
87.5%; Pred. No. 4e+02;
ative 0; Mismatches
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                                                                    294 GTGAAGGACCTGAGCC 309
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                                                                                                                                                                                        AAA30349 standard; DNA; 19
                                                                                                                                                                                                                                                         14-SEP-2000 (first entry)
Query Match
Best Local Similarity 87.5
Matches 14, Conservative
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22-OCT-1999;
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                                                                                                                                                                                                                            AAA30349;
                                                                                                                                                       RESULT 502
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The invention relates to a retinoid X receptor-gamma (RXR-gamma) knockout

mouse whose germ and somatic cells contain an insertion of an exogencus

mouse whose germ and somatic cells contain an insertion of an exogencus

containing portions of the RXR-gamma gene (exons 3 and 4) which

encode the entire DNA binding domain of RXR-gamma. The knockout mouse is

deficient in the normal expression of RXR-gamma. The invention

concompasses mice which are either homozygous or heterozygous for the

defective RXR-gamma gene, and also encompasses mammalian particularly

concontaining an exogenous DNA insert within exors 3 and 4. The

concontaining an exogenous DNA insert within exors 3 and 4. The

conclusion and antagonists of identifying RXR-gamma

agonists or antagonists using the transgenic mouse or marmalian cell

colline. The generically regimeered mouse and cell line are useful in

colline. The generically engineered mouse and cell line are useful in

collass of receptors. The mouse and cell line allow the investigation at

collass of receptors. The mouse and cell line allow the subscription of the importance of each of the RXR-gamma and in vivo levels of a system that lacks one or more

containing development and physiology. They are useful in studying any

contained development and physiology. They are useful in studying any

contained analysis of RXR-gamma reverse transcription-PCR (RT-PCR) primers used

contained analysis of RXR-gamma reverse transgenic mice of the invention. The

present sequence is an RT-PCR primer for exon B5 of the mouse RXR-gamma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New genetically engineered mice containing alterations in the gene encoding retinoid X receptor, useful for identifying agonists and antagonists of the receptors and in studying retinoic acid mediated gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
3.0%; Score 12.8; DB 1; Length 19;
Best Local Similarity 87.5%; Pred. No. 4e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19 BP; 4 A; 6 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 2; Col 12; 20pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    113 CCGCAGCAAGTACGGC 128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4 CCACAGCAAGTTCGGC 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 expression.
ð
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RESULT 503 AAA72197 ID AAA72197 standard; DNA; 19

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AAD19298;

RESULT 504

AAD19298/

WO200173035-A1

Mammalia Key allele 04-OCT-2001

Morahan G;

Page 244

rng.res

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Edinger S, Macdougall JR, Millet I, Ellerman K, Stone DJ;
Gerlach V, Grosse MM, Alsobrook JP, Lepley DM, Rieger D, Burgess CE;
Garana KJ, Spycek KA, Boldog FL, Li L, Padigaru M, Mishra V;
Patturajan M, Shenoy S, Rastelli L, Tchernev VT, Vernet CAM;
Zerhusen BD, Malyankar UM, Guo X, Miller CE, Gangolli EA;
hypertensive; haemostatic; cardiant; antianginal; dermatological; immunosuppressive; antiinflammatory; virucide; antibacterial; anti-HIV; antiparasitic; antiallergic; antiaschmatic; antirheumatic; antiarthritic; vulnerary; anorectic; antidiabetic; immunomodulator; antipsoriatic; nephotropro; kerolytic; antidiabetic; crebroprotective; antidiabetic; antidiatertility; antimanic; antidepressant; metabolic; cytostatic; tranquilizer; analgesic; probe; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention provides the protein and coding sequences of several novel human proteins, designated NOVX. These can be used in the treatment of, amongst others, cancers, autoimmune diseases, infections, inflammatory diseases, storage disorders, muscle disorders, neurodegenerative diseases and developmental defects. The present sequence is a PCR primer or probe used to isolate the sequences of the invention. All of the probes are modified at the 5' end by TBT and at the 3' end by TBMRA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cell culturing; embryonic stem; ES; central nervous system; fgf8; dopaminergic; cholinergic; serotonergic; antiparkinsonian; nootropic; neuroprotective; anticonvulsant; tranquilizer; vulnerary; neuroleptic; cerebroprotective; cell therapy; gene therapy; CNS; PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel isolated polypeptide, designated NOVX, useful for treating or preventing in NOVX-associated disorders e.g. cardiomyopathy, atherosclerosis, diabetes, cancer, allergy, asthma, Crohn's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
3.0%; Score 12.8; DB 1; Length 19;
Best Local Similarity 87.5%; Pred. No. 4e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Receptor fgf8 cDNA amplifying forward primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 211; 353pp; English.
                                                                                                                                                                                                                                                                                                                                                                                              29-NOV-2000; 2000US-0253834P.
30-NOV-2000; 2000US-0250926P.
25-JAN-22001; 2001US-0264180P.
20-AUG-2001; 2001US-0313656P.
05-OCT-2001; 2001US-0327456P.
28-NOV-2001; 2001US-00327456P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           70 ACTACGAGGGCCGCGC 85
                                                                                                                                                                                                                                                                                                                                                       29-NOV-2001; 2001WO-US048922.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-590741/63.
                                                                                                                                                                                                                                                            WO200257450-A2.
                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                           25-JUL-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The patent discloses a method of screening mammals for autoimmune diseases by identifying polymorphisms in interleukin (IL)-12 p40 gene. The methods and kits of the invention are used for screening individuals, families and populations for disease conditions or predispositions for the development of a disease condition which is characterised, exacerbated or associated with Th1/Th2 dysregulation in amammal. They are used to treat, prevent or disapnose autoimmune diseases such as IDDM (Insulin dependant diabetes mellitus). The present DNA sequence is mammalian IL-12 p40 intron 7 allelic variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; NOVX; autoimmune disease; cancer; infection; inflammatory disease; storage disorder; muscle disorder; neurodegenerative disorder; nootropic; developmental defect; neuroprotective; antiparkinsonian; hypotensive;
                                                                                                                                                                                                                                                         Interleukin-12; IL-12 p40; autoimmune disease; Th1/Th2 dysregulation; therapy; allelic variant; insulin dependant diabetes mellitus; IDDM; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Screening mammals for autoimmune diseases such as diabetes, comprises identifying polymorphisms in interleukin ({\rm IL})-12 p40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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3.0%; Score 12.8; DB 1; Length 19;
Best Local Similarity 87.5%; Pred. No. 4e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19 BP; 5 A; 4 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                Mammalian IL-12 p40 intron 7 allelic variant #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (HALL-) HALL INST MEDICAL RES WALTER & ELIZA.
                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
replace(10, A)
/*tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 21; Page 42; 115pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                118 GCAAGTACGGCATGCT 133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABT06307 standard; DNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27-MAR-2001; 2001WO-AU000340
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-MAR-2000; 2000AU-00006466
15-MAY-2000; 2000US-0204366P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GCAAGTCCGCCATGCT 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                    1298/c
AAD19298 standard; DNA; 19
                                                                                                                                                                      18-DEC-2001 (first entry)
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24-OCT-2002

ABT06307/ ID ABT0 XX AC ABT0 XX DT 24-0 DT 24-0 XX Huma XX Huma XW Huma XW Geve

ABT06307;

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Wed Apr 21 12:58:21 2004
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01-MAY-2001; 2001WO-US014051
   WO200183715-A2.
 Homo sapiens
     08-NOV-2001
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01-MAY-2000; 2000US-0201005P (USGO ) US GOVERNMENT. (LDES/) LEE S. (LUME) LUMELSKY N. (STUD/) STUDER L. (MCKA/) MCKAY R D G.

Studer L, Mckay RDG Lumelsky N, Lee S,

WPI; 2002-049345/06.

Culturing cells such as neuronal cells for use in treating neurological disorders, comprises generating embryoid bodies from undifferentiated embryonic stem cells, selecting precursor cells, expanding and differentiating them.

Example 10; Page 41; 66pp; English.

The invention provides a method of culturing cells. The method involves expanding a culture of undifferentiated embryonic stem (ES) cells, contrain method embryonic by culturing the bodies (EB), culturing the bodies to select for central nervous system (CNS) precursor cells (PC), culturing policy of the expanded PC to form a culture of differentiated neuronal cells are useful for culturing the expanded PC to form a culture of differentiated neuronal cells which are useful for treating and cells. The method is useful for culturing undifferentiated neuronal cells which are useful for treating and neurological disorder, especially Parkinson's disease in a patient. A gene product such as tyrosine hydroxylase, nerve growth factor (NGF), brain derived neuronal cells which are useful for culturing dopaminergic, cholinergic actor (GDNF) NT-3, and NT-4/5 can be introduced into a brain of a subject. The method is useful for culturing dopaminergic, cholinergic and serotonergic neuronal cells. The differentiated neuronal cells are useful for treating neurological disorders such as Huntington's disease, all planer's disease, multiple sclerosis, severe seizure disorders (contrained policy, familial dysautonomia as well as injury or trauma to the nervous system such as neurotoxic injury or disorders for measure disorders cuch as addiction and schizohrenia, cerebrovascular disorders such as stroke and CNS disorders resulting from adjing. Assays are useful for developing drugs capable of regulating the survival, proliferation or genesis of neuronal cells and to screen for antagonist or agonist of companies 20-40% dopaminergic neurons and 1-3% astrocytes are useful for studying the mechanism of neuronsmitter synthesis and release, particularly for serotonin and dopamine, neuronal cells survival, and the companies 20-40% dopaminergic neurons and 1-3% astrocytes are useful for studying the mechanism of neuroransmitter synthesis and release, particularly for serotonin and dopaminer neuronal cells survival, and the electrophysiochem

Sequence 19 BP; 5 A; 5 C; 7 G; 2 T; 0 U; 0 Other;

Gapa ö Query Match
3.0%; Score 12.8; DB 1; Length 19;
Best Local Similarity 87.5%; Pred. No. 4e+02;
Matches 14; Conservative 0; Mismatches 2; Indels

à g RESULT 507 ABS97846 ID ABS9784

ABS97846 standard; DNA; 19 BP.

(first entry) 23-DEC-2002

sulfotransferase thermolabile (STM) gene PCR primer #1.

Human; ss; primer; cytochrome P450 A1; CYP4501A1; UGT2B4; MDR1; PCR;

KW cytochrome P450 A2; CYP4501A2; cytochrome P450 02E; CYP45002E1; LTF;

KW adrenergic receptor beta1; ADBR1, aryl hydrocarbon; AHR; MRP3; NRT12;

KW adrenergic receptor nuclear translocator; ARNT; cathepsin S; CTSS;

KW aryl hydrocarbon receptor nuclear translocator; ARNT; cathepsin S; CTSS;

KW cyclooxgenase 2; EPRX2; 5-lipoxygenase activating protein; FLAP;

KW glutathione-S-transferase 12; GST12; histamine-N-methyl transferase;

KW MDPH quinone oxidoreductase 2; NQO2; sulfotransferase thermolabile; STM;

WDP-gluturonosyl transferase; UGT2B15; urokinase receptor;

W UDP-gluturonosyl transferase; UGT2B15; urokinase receptor;

W multidrug resistance 1; lactotransferrin; orphan nuclear receptor;

W multidrug resistance associated protein 3; cancer; prostate;

R multidrug resistance associated protein 3; cancer; prostate;

R multidrug resistance succeptor; CHWR1; CHWR2;

R multidrug resistance wasciated protein 3; cancer; prostate;

R multidrug resistance succeptor; CHWR1; CHWR2;

R multidrug resistance wasciated protein 3; cancer; prostate;

R multidrug resistance succeptor;

R multidrug resistance wasciated protein 3; cancer; prostate;

R multidrug resistance wasciated protein; cHWR2; CHWR3;

R multidrug resistance system; pulmonary; lumunological.

WO200257410-A2.

25-JUL-2002.

28-NOV-2001; 2001WO-US044838.

28-NOV-2000; 2000US-00724389

(DNAS-) DNA SCI LAB INC.

Guida M, Hall J;

WPI; 2002-698522/75.

Isolated nucleic acid molecules having polymorphisms in known human genes eg. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for disorder-related tratts.

Example 17; Page 131; 714pp; English.

This invention relates to the sequence of an isolated mucleic acid
molecule comprising at least one base variation from that of a known
to wolecule comprising at least one base variation from that of a known
to wrotcohrome P450 al (CYP4501A1), Tytochrome P450 A2 (CYP4501A2),
cytochrome P450 02B1 (CYP45002B1), adrenergic receptor Detail (ADBR1),
aryl hydrocarbon (AHR), aryl hydrocarbon receptor nuclear translocator
(ARMY), cathepsin S (CTSS), Oyclooxgenase 2 (CMCX), diazegam bhiding
inhibitor (DBI), epoxide hydroxylaes 2 (EHKZ), 5-lipoxygenase activating
cinhibitor (DBI), poxide hydroxylaes 2 (EHKZ), bistamine-N-methyl
transferase (NNMY), NADPH quinone oxidoreductase 2 (NGO2),
ctransferase thermolabile (STM), UD-glucuronosyl transferase 284
(UGT2B1,) unorplusinencyl transferase 287 (UGT2B1), UDP-glucuronosyl
transferase (UGT2B15), urokinase receptor (NPA), multidrug resistance 1
(MDRI), lactotransferrin (LTF), multidrug resistance associated protein 3
(MDRI), lactotransferrin (LTF), multidrug resistance associated protein 3
(MDRI), lactotransferrin (LTF), multidrug resistance associated protein 3
(CTRR2), orphan nuclear receptor (NRI2), or acceptor(LNA) exceptor 1, 2, 3, 4, or 5 (CHMR), CHMR2, CHMR3, CHMR4, or CHMR5) sequence
creeptor 1, 2, 3, 4, or 5 (CHMR), CHMR3, CHMR4, or CHMR5) sequence
creeptor 1, 2, 3, 4, or 5 (CHMR), cHMR3, CHMR4, or CHMR5) sequence
creeptor 1, 10x489 markers for locating and characterising the genes that
care responsible for specific traits within the genome and eventually
care responsible for specific traits within the genome and eventually
care responsible end septonsible for a variety of disorder-related
complexession, mutation or underexpression, which may be used in diagnosing
complexession, mutation or underexpression, which may be used in diagnosing
complexession, mutation or underexpression, mutation are useful for screening individuals for altered drug

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metabolism. The polymorphic sequences contained in CYP4501A1, CYPP4501A2, AMR, WDR1 and/or MDR3 may also be used to screen individuals for susceptibility to cancer. Polymorphic sequences in ADRB1 or CHMR2 are used to screen for altered cardiovascular function, in CXX2 for altered susceptibility to colorectal tumours, in DB1 or CHMR1 for altered central nervous system function, in FLAP and HNMT for altered pulmonary, immunological or haematological function, in KLK2 for altered serine protease activity in the prostate, in LTF for altered immunological or haematological function, in CHMR3, CHMR4 or CHMR5 for altered central and peripheral nervous system function. The present sequence represents a PCR primer used to amplify the sequences of the invention
          8888888888888888
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Sequence 19 BP; 4 A; 3 C; 7 G; 5 T; 0 U; 0 Other;

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3.0%; Score 12.8; DB 1; Length 19;
87.5%; Pred. No. 4e+02;
iive 0; Mismatches 2; Indels 0; Gaps
                 1 Similarity 87.5
 Query Match
                     Local
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55 CAGAGGAGTCTCTGCA 70 2 CAGAGGAGTTTCTTCA 17 ð

ABZ69849 standard; RNA; 19 BP (revised)
(first entry) 27-OCT-2003 10-APR-2003 ABZ69849; ABZ69849/c 

HIV-1 strain HXB2 RNAi target sequence 2.

Ribozyme; Rz2; pharmaceutical carrier; haematopoietic; anti-HIV; virucide; cytostatic; antianaemic; cardiant; gene therapy; cell therapy; antianse therapy; HV; haemoglobinopathy; leukocyte; Fanconi's anaemia; chronic granulomatous disease; Gaucher's disease; GGFD deficiency; cardiovascular disease; HIV-1-HXB2; ss.

Human immunodeficiency virus 1.

WO2003006691-A1.

23-JAN-2003

10-JUL-2002; 2002WO-US021907. 10-JUL-2001; 2001US-0304127P.

21-DEC-2001; 2001US-0343484P. 04-JUN-2002; 2002US-0386063P.

(GENE-) GENE SHEARS PTY LID.

Macpherson J, Fanning G, Gerlach W; Sun L, Amado R, WPI; 2003-221763/21. Symonds GP,

New composition comprising CD34 hematopoietic cells transduced with a viral construct expressing an anti-HIV agent, useful for treating HIV, AIDS, and diseases of the blood and immune systems, e.g. Fanconi's anemia or cancer,

Example 5; Page 112; 157pp; English.

The invention relates to a novel composition comprising a pharmaceutical carrier and haemacopoietic dells transduced with a viral construct expressing an anti-thy agent. A composition of the invention has virucide, cytostatic, antianaemic, anti-HIV, and cardiant activity. The compositions may have a use in gene therapy, cell therapy, and antisense therapy. The composition is useful in the manufacture of a medicament for the treating HIV. The composition is useful in the annufacture of a medicament for

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variety of diseases in which there is a genetic aspect, such as diseases of the blood and immune systems, including haemoglobinopathies, defects of leukocyte production or function including cancers, AIDS/HIV, viral infections, Iysosomal storage diseases and stem cell defects such as Panconi's anaemia, chronic granulomatous diseases, Gaucher's disease, GeFD deficiency, and cardiovascular diseases. The present sequence represents a highly conserved RNMi target sequence from HIV-1 HXB2. (Updated on 27-OCT-2003 to standardise OS field)
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Seguence 19 BP; 5 A; 6 C; 5 G; 3 T; 0 U; 0 Other;

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3.0%; Score 12.8; DB 1; Length 19; 87.5%; Pred. No. 4e+02; tive 0; Mismatches 2; Indels
                                      14; Conservative
                   Local Similarity
   Query Match
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ACF35801,

ACF35801 standard; DNA; 19 BP.

06-NOV-2003 (first entry)

Human GPR43 receptor DNA amplifying gene-specific forward primer.

GPR41; G-protein coupled receptor; fatty acid; antiemetic; antimigraine; neuroleptic; antidepressant; tranquilliser; neuroprotective; noctropic; antiparkinsonian; anticonvulsant; antimanic; antigatic; cytostatic; metabolic; immunomodulator; antiathmatic; cardiant; hypotensive; osteopathic; antianginal; antiuloer; antiallergic; cerebroprotective; human; RT-PCR; primer; ss.

Homo gapiens.

WO2003057730-A1.

06-JAN-2003; 2003WO-EP000042.

07-JAN-2002; 2002US-0346396P 

(EURO-) EUROSCREEN SA.

Brezillon S, Lannoy V, Parmentier M; Le Poul E, Detheux M,

WPI; 2003-598359/56.

Identifying agent that modulates GPR43 function, useful for treating migratine, soltzophrenia, anxiety, by measuring binding of GPR43 polypeptide to short chain fatty acid in presence and absence of candidate modulator.

Example 2; Page 80; 136pp; English.

The invention relates to identifying an agent that modulates function of G-protein coupled receptor GPR43. The method involves measuring the binding of GPR43 polypeptide to short chain fatty acid [II] in presence and absence of [III], or measuring signaling activity of GPR43 contacted with [II] in presence and absence of [III], or measuring signaling activity of GPR43 in presence of [II] and comparing the activity to activity measured in a sample in which GPR43 is contacted with [II] at its ECSO. The agents identified are useful for modulating the activity of GPR43 in a call and for modulating polencyhounclear (PMN) chemotaxis in a mammal. The agents are useful for manufacture of medicament for tracting GPR43-related diseases or PMN chemotaxis-related diseases or disorders such as vomiting, migraine, schizophrenia, manic depression, anxiety, dementia, neurodegenerative diseases such as

Alzheimer's disease and Parkinson's diseases and dyskinesias, such as bulbutington's disease. They are also useful for preventing, improving or correcting dysfunction or diseases e.g., pain, cancer, anorexia, bulimia, asthma, acute heart failure, hypertension, osteopoxosis, urinary tetention, angina pectoris, myocardial infarction, ulcers, allergies, stroke, and schizophrenia. The present sequence represents a GPR43 genespecific primer used in semi-quantitative RT-PCR reactions

88888888888

Sequence 19 BP; 2 A; 7 C; 5 G; 5 T; 0 U; 0 Other;

0; Gaps Query Match 3.0%; Score 12.8; DB 1; Length 19; Best Local Similarity 87.5%; Pred. No. 4e+02; Matches 14; Conservative 0; Mismatches 2; Indels

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210 RESULT

ADE65585 standard; RNA; 19 BP. ADE65585

ADE65585;

29-JAN-2004 (first entry)

Human c-fos transcript target sequence/siNA upper strand, SEQ ID NO:40.

RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairbin RNA; shrah; expression modulation; gene therapy; drug soreening; diagnosis; therapentic target identification; pharmacogenomics; gene function analysis; gene mapping; central nervous system disorder; Alzheimer's disease; palepsy; dementia; parkinson; disease; mintington's disease; epilepsy; dementia; amyotrophic lateral sclerosis; cancer; proliferative disease; restenosis; polycystic kidney disease; inflammatory disease; transplant rejection; viral infection; HIV infection; autoimmune disease; transplant rejection; vasoricopic; antibalkinsonian; neuroprotective; cytostatic; antiinflammatory; antiallergic; virucide; anti-HIV; immunosuppressive; anticonvulsant; nephrotropic; human; c-fos; target sequence; ss.

Homo sapiens.

WO2003070914-A2.

28-AUG-2003.

20-FEB-2003; 2003WO-US005162.

20-FEB-2002; 2002US-0358580P. 11.MAR-2002; 2002US-0363124P. 06-UUN-2002; 2002US-0386782P. 29-AUG-2002; 2002US-0406378P. 05-SEP-2002; 2002US-0408378P. 

(SIRN-) SIRNA THERAPEUTICS INC.

09-SEP-2002; 2002US-0409293P. 15-JAN-2003; 2003US-0440129P.

Mcswiggen J, Beigelman L;

WPI; 2003-679877/64.

New short interfering nucleic acid downregulates expression of the c-fos gene useful for treatment and diagnosis of diseases, e.g. cancer and inflammation.

Example 3; SEQ ID NO 40; 145pp; English.

The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the human c-fos gene by RNA interference. The

cc sinka may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the sinka include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The sinka can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised (expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo dealivery of siNA, conjugates and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the c-fos gene in cells, tissue explants or organisms of siNA, by ex vivo gene therapy), or in grafts and transplants for the creatment of a variety of conditions. They may be used for treating central nervous system lesions and injuries (e.g., Alzheimer's disease, elements of anyotrophic lateral sclerosis), various cancers other proliferative diseases (e.g., restenosis and polycystic kidney disease); inflammatory condourned diseases; varial infections (including HIV infection); autoimmune diseases; varial infections (including gene contrian), and gene mapping (e.g., of single mucleotide polymorphisms).

The present sequence represents the upper strand of a human c-fostarded double-stranded siNA, which is identical to the c-fos transcript target sequence.

%XGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG

Sequence 19 BP; 6 A; 5 C; 6 G; 0 T; 2 U; 0 Other;

. 0 3.0%; Score 12.8; DB 1; Length 19; 75.0%; Pred. No. 4e+02; ative 2; Mismatches 2; Indels Local Similarity 75.0 les 12; Conservative Query Match Best Loca Matches

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Bb. ADE65701 Standard; RNA; 19 29-JAN-2004 (first entry) ADE65701; RESULT 511 ADE65701/c

Human c-fos siNA lower strand, SEQ ID NO:156.

RNA interference; short interfering nucleic acid; siNA; shNA; shNA; shNA; shNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; central nervous system disorder; Alzheimer's disease; parlings; parkingon's disease; proliferative dimentia; amyotrophic lateral sclerosis; cancer; proliferative disease; restenosis; polycystic kidney disease; inflammatory disease; transplant rejection; viral infection; HIV infection; autoimmune disease; transplant rejection; vasorropic; nootropic; antiparkinsonian; neuroprotective; cytostatic; antiinflammatory; antiallergic; virucide; anti-HIV; immunosuppressive; anticonvulsant; nephrotropic; human; c-fos; ss. 

Homo gapiens.

WO2003070914-A2.

28-AUG-2003.

20-FEB-2003; 2003WO-US005162

20-FEB-2002; 2002US-0358580P. 11-MAR-2002; 2002US-0363124P. 06-UJN-2002; 2002US-0386782P. 29-AUG-2002; 2002US-0406784P.

The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the human c-fos gene by RNA interference. The siNAS may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense coligonucleotide. Specifically, the siNAS include short interfering RNA (siRNA), double-stranded RNA, mitto-RNA include short interfering RNA (siRNA). The siNAS can be unmodified or chemically modified, can contain covertor or enzymatically synthesised. The invention also relates to kits or the in vitro or in vivo delivery of siNA; conjugates and/or complexes of six, and vectors that express siNA. The siNAS are used to modulate expression of the c-fos gene in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating central nervous system lesions and injuries (e.g., Alzheimer's disease, Parkinson's disease, Huntington's disease, epilepsy, dementia or amyotrophic lateral sclerosis, various cancers; other proliferative diseases (e.g., restenosis and polycystic kidney disease); inflammatory and/or allergic diseases; wiral infections (including HIV inflection); autoimmume diseases; and transplant rejection. The sinAs are also useful function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human c-fos-New short interfering nucleic acid downregulates expression of the c-fos gene useful for treatment and diagnosis of diseases, e.g. cancer and Example 3; SEQ ID NO 156; 145pp; English. (SIRN-) SIRNA THERAPEUTICS INC. 09-SEP-2002; 2002US-0409293P. 15-JAN-2003; 2003US-0440129P. WPI; 2003-679877/64. Mcswiggen J, inflammation. 

Sequence 19 BP; 2 A; 6 C; 5 G; 0 T; 6 U; 0 Other;

Gaps ö Match 3.0%; Score 12.8; DB 1; Length 19; Local Similarity 87.5%; Pred. No. 40+02; Local Similarity 0; Mismatches 2; Indels 14; Conservative 0; Mismatches 2; Indels 14; Conservative Query Match Best Loca Matches

CCAAGCTGGTGAAGGA 301 CCAACCTGCTGAAGGA 3 18

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ABZ86355 standard; DNA; 20 BP ABZ86355; RESULT 512 ABZ863 

(first entry) 17-OCT-2003

Human oligonucleotide sequence

Human; antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

Pharmaceutical composition for treating ailments associated with impaired respiration, has Oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or WPI; 2003-229219/22.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entiting amount of the invention of the antiinflammatory servoid and ubiquinone. A composition of the invention of mas antiinflammatory attiallargic, antiasthmatic, hypotensive, commonsuppressive, and cytostatic activity. The composition may have a immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory lung or malignam is useful for treating or conforming the prophylactic or therapeutic respiratory effect of an or infilammatory steroid in a subject, for reducing levels of adenosine or receptor, producing bronchodilation, increasing levels of adenosine creeptor, producing bronchodilation, increasing levels of wbiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung alleraties, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at the printed or 

Sequence 20 BP; 5 A; 6 C; 6 G; 3 T; 0 U; 0 Other;

Gaps ö Query Match
3.0%; Score 12.8; DB 1; Length 20;
Best Local Similarity 87.5%; Pred. No. 4.5e+02;
Matches 14; Conservative 0; Mismatches 2; Indels

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16 Argreakcrecreer 1 ઠે 셤

AAT06919 standard; DNA; 19 BP. 04-JUL-1996 (first entry) AAT06919; AAT06919 BXBXSXXXXXXXXXXXXXXXXX

RESULT 513

Chromosomal locus E17 primer #1.

prostate/colon tumour suppressor gene; allelic loss; colorectal cancers; microsatellite analysis; sequence tagged site; primer; probe; PCR; amplification; chromosome; ss.

WO9532214-A1

30-NOV-1995

WO200285308-A2

WEG APT 41 14.30.41 400

31-OCT-2002.

Beigelman L;

(EPIG-) EPIGENESIS PHARM INC.

Katz E, Pabalan J, Aguilar D; Li Y, Sandrasagra A, K. Tang L, Shahabuddin S; Nyce JW, | Miller S,

ubiquinone.

Claim 15; SEQ ID NO 1597; 872pp; English.

190 ATATCCACTGCTCGGT 205

Synthetic.

Disclosure; Page 35; 122pp; English.

95WO-US006593 94US-00246604

Isaacs WB;

INC.

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Primers AAT06887-932 were used to analyse the breakpoints at chromosomal locus 8p2-21, contained in patients having prostate cancer, by microsatellite analysis and sequence taggaed sites (STS). The region contains a prostate/colon tunour suppressor gene (PTS). The primers and amplified fragments were used to screen a YAC library of prostate cancer DNA to isolate the PSTG (AAT06880), which can be used in the diagnosis and treatment of prostate and colorectal cancers. The primers AAT06919-20 amplify a 121 bp fragment from chromosomal locus BI?
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New nucleic acid encoding mammalian tub protein - useful for diagnosis and treatment of body wt. disorders, esp. obesity, and for screening for
                                                                                                                                      'DNA encoding a prostate tumour suppressor protein - from chromosome for the diagnosis and treatment of prostatic and colorectal cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   tubby; tub; CBT9 gene; body weight; obesity; cachexia; anorexia;
                                                                                                                                                                                                                                                                                                                        Sequence 19 BP; 7 A; 6 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                             Disclosure; Page 86; 122pp; English
                                                                                                                                                                                                                                                                                                                                                                                                     182 CAAGGCACATATCCACTGC 200
                                                                                                                                                                                                                                                                                                                                                                                                                       CAAGGCATATCACAACTGC 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (MILL-) MILLENNIUM PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAT48575 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human tub gene primer R12
                                                                                                                                                                                                                                                                                                                                                                             15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1997-108751/10.
                                                                                                             WPI; 1996-020526/02.
                                                                                                                                                                                                                                                                                                                                                                Sest Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  disorders; ss.
                                                           (CANJ-) CANJI
                                                                                      Bookstein R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9702048-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28-JUN-1996;
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28-SEP-1995;
09-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19-0CT-1997
            22-MAY-1995;
                                    20-MAY-1994;
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20-JUL-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
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                                                                                                                                         New DNA
                                                                                                                                                                                                                                                                                                                                                                             Matches
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AAT48575/
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BP

(first entry)

95US-000604P. 95US-0001273P. 95US-0001444R. 95US-0002759P. 95US-0004424P.

96US-00631200

96WO-US011186

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nucleic acid sequence determination - comprising synthesising chain extension products, which are indicative of positions of selected species of nucleotide in nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This sequence represents a primer for exon 3 of the HLA-C gene. This sequence can be used in the method of the invention for determining the position of at least one selected species of nuclectide, in a region of interest, in a sample. The method comprises combining the sample with a reaction mixture to synthesise chain extension products indicative of the positions of the species of nuclectide in the region of interest and evaluating the products produced, characterised in that the sample, which is combined with the reaction mixture, and contains target and non-target nucleic acid polymers in natural abundance. The method can be used to detect mutations, particularly mutations of medical significance, in samples
                                                  The murine and human tub gene (AAT48550 and AAT48551 respectively) products are wild-type, expressed in the hypothalamus. The form lacking exon 5 is produced by alternative splicing. The products participate in the control of mammalian body weight. Messuring tub expression and detection of tub gene mutation are used to diagnose body weight disorders, esp. obesity, cachexia and anorexia, or related sensory and fertility defects
                                                                                                                                                                                                                                              Gape
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  detection;
human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Shipman
                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                Score 12.6; DB 1; Length 19;
Pred. No. 4.4e+02;
                                                                                                                                                                                                                                                4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR primer; amplify; pathogen identification; mutation uncleic acid analysis; microorganism characterisation; HLA type determination; HLA-C gene exon 3; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lacroix J,
                                                                                                                                                                                Sequence 19 BP; 6 A; 6 C; 7 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5' vgicwsp5 primer for exon 3 of HLA-C gene.
                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Larson MT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 6; Page 24; 69pp; English.
                                                                                                                                                                                                                                                                              132 CIGGCCCCCCTGGCGGTGG 150
                                                                                                                                                                                                                                                                                                            19 chracchachachan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          96US-00640672.
96US-00684498.
97US-00807138.
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                                                                                                                                                                                                                                                                                                                                                                                           BB
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                                                                                                                                                                                                                3.0%;
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                                                                                                                                                                                                                                                                                                                                                                                           AAT99886 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                               Local Similarity 78.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1997-549755/50.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-MAY-1996;
19-JUL-1996;
27-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Leushner J,
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                                                                                                                                                                                                                                                                                                                                                                                                                           AAT99886;
                                                                                                                                                                                                                   Query Match
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Gaps

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3.0%; Score 12.6; DB 1; Length 19; 78.9%; Pred. No. 4.4e+02; ive 0; Mismatches 4; Indels

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The present primer was used in the mapping of a gene encoding 2 forms of a prostate/colon tumour suppressor (P/CTS). The P/CTS gene was isolated by analysis of allelic loss in patients with prostate cancer, and was putatively located to the chromosomal location 8922-21 via microsatellite analysis and the use of sequence tagged sites (STS). Primers and probes derived from the gene can be used to screen lambda cDNA libraries for genes encoding P/CTS form 1 and 2. The P/CTS or its cDNA can be used in the diagnosis and treatment of prostate and colorectal cancers. (Updated on 25-WAR-2003 to correct PA field.) (Updated on 25-WAR-2003 to correct
                                                                                                                                                                         ..
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Prostate/colon tumour suppressor; allelic loss; prostate cancer; colorectal cancer; microsatellite analysis; sequence tagged site; STS; amplification; chromosomal location 8q22-21; probe; primer; gene mapping;
derived from a human patient, animal, plant or microorganism, determine HLA type ancillary to transplant procedures, detect and identify microorganisms, particularly pathogenic microorganisms, in a sample and in in situ sequencing reactions to produce sequencing fragments in a histological specimen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New prostate/colon tumour suppressor gene - mapped to a locus on human
                                                                                                                                                                       0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Primer E17 for mapping prostate/colon tumour suppressor gene.
                                                                                                                                  / Match 3.0%; Score 12.6; DB 1; Length 19; Local Similarity 78.9%; Pred. No. 4.4e+02; nes 15; Conservative 0; Mismatches 4; Indels
                                                                                                   G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 7 A; 6 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 26; 45pp; Japanese.
                                                                                                                                                                                                          378 GACCGCGACGACGCGCCCA 396
                                                                                                                                                                                                                                           gacciedeseseseseses 19
                                                                                                                                                                                                                                                                                                                                BP
                                                                                                     Sequence 19 BP; 2 A; 6 C; 11
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(UYJO ) UNIV JOHNS HOPKINS.
                                                                                                                                                                                                                                                                                                                                AAT64713 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                    (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            diagnosis; treatment; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Isaacs WB, Bookstein R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1997-275447/25.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-MAY-1995;
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12-FEB-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-FEB-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              chromosome 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-APR-1997.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                   AAT64713;
                                                                                                                                                                                                                                           -
                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                 RESULT 516
                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                   AAT64713
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This sequence represents a PCR primer for the human ACE (angiotensin converting enzyme) gene, and can be used in the method of the invention.

C determining the sequence of at least one polymorphic site in the ACE (angiotensining the sequence of at least one polymorphic site in the ACE (angiotensin converting enzyme), ACT (angiotensinogen) and/or ATI (type 1 converting enzyme), ACT (angiotensinogen) and/or ATI (type 1 angiotensin II receptor) genes, and comparing the polymorphic pattern of with that in patients with predetermined markers of Status. The method is used to assess blood pressure or electrocardiographic profile, to diagnose a cardiac condition such as (silent) myocardial infarction (MI), consequence or the certocardiographic profile, to response to treatments with ACE inhibitors, angiotensin II receptor antagonists, diuretics, alpha- or beta-adrenergic receptor antagonists, etc. It is also used to identify susceptibility to cardiovascular confises. Libraries of tucleic acids contraining polymorphic positions in the genes are used to screen for cardiovascular agents. The nucleic acids contained in the library can be is used as source of probes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Assessing cardiovascular status in humans by polymorphic analysis - of genes for angiotensin converting enzyme, angiotensinogen and angiotensin II receptor, used to diagnose predisposition to disease and to predict effect of therapy.
                                                                                                                                                                                                                                                                    PCR primer; human; ACB; angiotensin converting enzyme; angiotensinogen; cardiovascular status; AGT; ATI; type 1 angiotensin II receptor; stroke; polymorphic pattern; Blood pressure; electrocardiographic profile; cardiac condition diagnosis; myocardial infarction; atherosclerosis; hypertension; cardiovascular disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 19 BP; 7 A; 3 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Norberg LT, Andersson MK, Lindstroem PHR;
                                                                                                                                                                                                                                           Primer ACE/82RB for human ACE gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Page 27; 71pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  360 GACTICCICACTITCCIGG 378
182 CAAGGCACATATCCACTGC 200
                              1 CAAGGCATATCACAACTGC 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GATTICTICACCICCIGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98WO-IB000475.
                                                                                                                                 AAV08577 standard; DNA; 19
                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (EURO-) EURONA MEDICAL AB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1998-568361/48.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      04-APR-1997;
                                                                                                                                                                                                         15-FEB-1999
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Best Local Si
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
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                                                                                                                                                                 AAV08577;
                                                                                          RESULT 51
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Gaps

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Query Match
3.0%; Score 12.6; DB 1; Length 19;
Best Local Similarity 78.9%; Pred. No. 4.4e+02;
Matches 15; Conservative 0; Mismatches 4; Indels

WO9920749-A1

Synthetic

AAX16754;

RESULT 518

AAX16754,

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The specification describes a method for screening for functional polypeptides which bind a ligand. The method comprises contacting a repertoire of polypeptides with a generic ligand, and then screening selected functional polypeptides with a target ligand. The method permits the removal from a chosen repertoire of polypeptides, those which are non-functional, e.g. as a result of the introduction of frame-shift be or are incapable of binding untants or expression mutants which would be or are incapable of binding to any target ligand. The method also polypeptides which are functional, well folded and highly expressed. The polypeptides obtained can be used in diagnostic, prophylactic and therapeutic procedures. PCR primers AAX35946-48 were used to amplify a germiline V gene fragment, which was used in the construction of libraries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PG1 gene; biallelic marker; PCR primer; PG1-related biallelic marker; cancer; prostate cancer; diagnosis; therapy; prostate specific antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.0%; Score 12.6; DB 1; Length 19; 78.9%; Pred. No. 4.4e+02; tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                        Screening for functional polypeptides which bind a ligand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19 BP; 2 A; 7 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PCR primer for PG1 biallelic marker 99-123-184.
                                                                                                                                                                                                                                                                                                                                                  Example 2; Page 49; 67pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         270 CIGGAGCAGGGGGCACCA 288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19 CreakGCCTGGCGGACCCA 1
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98US-0099658P.
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                                                                         98WO-GB003135
                                                                                                                97GB-00022131
                                                                                                                                   97US-0065428P.
97US-0066729P.
                                                                                                                                                                                          (MEDI-) MEDICAL RES COUNCIL.
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                                                                                                                                                                                                                                   Tomlingon I, Winter G;
                                                                                                                                                                                                                                                                       WPI; 1999-288302/24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PSA; human; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-JUL-1999.
                                                                                                                                   13-NOV-1997;
21-NOV-1997;
                                                                         20-OCT-1998;
                                                                                                                20-OCT-1997;
                                   29-APR-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Primers AXX16733-X16754 are examples of primers which can be used to PCR amplify the human "tub" gene (AAX16702) exons. The invention relates to a method for identifying compounds that modulate tub protein activity, especially its interaction with proteins containing an SH2 domain. The method can be used for identifying compounds which modulate tub protein activity for use in the treatment of mammalian body weight disorders including obesity, cachexia and anorexia
                                                                                                                                                                                                                 Mouse, wild type, tubby, identification, SH2 domain, mammal, obesity, body weight disorder, cachexia, anorexia, primer, PCR, amplification, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Identifying compounds which modulate tub protein activity - by detecting compounds which alter the interaction of tub protein with a SH2 containing peptide, used to develop agents for treating e.g. obesity,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Screening, functional polypeptide, ligand, non-functional, enrichment, single chain antibody, PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5' primer used to amplify germline V gene segment DP-47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 19 BP; 6 A; 6 C; 7 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                             Human tub gene exon 12 R12 primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Col 22; 95pp; English
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97US-00829553.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (MILL-) MILLENNIUM PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAX35946 standard; DNA; 19
                                                            AAX16754 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-JUL-1999 (first entry)
                                                                                                                                          27-APR-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kapeller R, Moore KJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cachexia or anorexia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1999-130383/11.
                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                   02-SEP-1997;
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Synthetic

AAX35946;

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519

RESULT

AAX35946,

Query Match

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Gaps ö

Page

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The invention relates to a mammalian PG1 gene and protein, and a set of PG1 biallelic markers. The PG1 polynuclecide and biallelic markers are used in a hybridisation assay, a sequencing assay, or in an allelespecific amplification assay for determining the identity of a nucleotide at a PG1-related biallelic marker. The methods can be used to detect and to assess the risk of developing cancer or prostate cancer. Early-stage diagnosis of prostate cancer relies on prostate specific antigen (PSA) chapter, the effectiveness of this is limited due to its inability to discriminate between malignant and non-malignant affections of the organ. A need exists for both a reliable diagnostic procedure curative treatments of the disease. The PG1 gene can be used for detection of prostate cancer, and the risk of developing it in the future, and can also be used to determine therapies for the disease
                                                                                                       Use of a prostate cancer associated gene and biallelic markers derived
                      Bougueleret L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 19 BP; 6 A; 8 C; 2 G; 3 T; 0 U; 0 Other;
                   Chumakov I,
                                                                                                                                                                              Claim 4; Page 367; 385pp; English.
                   Cohen D, Blumenfeld M,
                                                                  WPI; 1999-405178/34
                                                                                                                                         from it.
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Query Match
3.0%; Score 12.6; DB 1; Length 19;
Best Local Similarity 78.9%; Pred. No. 4.4e+02;
Matches 15; Conservative 0; Mismatches 4; Indels 289 AGCTGGTGAAGGACCTGAG 307 19 AGCTGGTGAATGTTCTGGG 1 셤 ò

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Gaps

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Human HPC2 cDNA exon 24 3'UTR mutation screening primer SEQ ID NO: 185 AAA60364 standard; DNA; 19 (first entry) 07-DEC-2000 AAA60364; AAA60364 RESULT 

Human, mouse, prostate cancer predisposing gene, HPC2; human chromosome 17p, gene therapy, peptide therapy, drug design, PCR primer, sequencing primer; ss.

New hairpin and hammerhead ribozyme for inhibiting restenosis, cleaves RNA encoding a cyclin or cell-cycle dependent kinase other than CDK1, PCNA and Cyclin B1.

Disclosure; Page 52; 109pp; English.

Robbins JM;

Barber JR,

Tritz R, Welch PJ, WPI; 2000-412314/35.

domo sapiens

18-MAY-2000

40200027864-A1.

05-NOV-1999;

98US-0107468P 06-NOV-1998;

(MYRI-) MYRIAD GENETICS INC.

Pavtigian SV, Teng DHF, Simard J, Rommens JM; WPI; 2000-376481/32.

Example 5; Page 62; 157pp; English.

The present sequence is a primer used in the isolation of the human and murine prostate cancer predisposing genes HPC2 and Mm.HPC2. The human

Human prostate cancer (HPC)2 nucleic acids, polypeptides, and antibodies, useful for treatment and diagnosis of prostate cancer.

523

RESULT

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version of the gene is found on chromosome 17p. Some alleles cause a predisposition to cancer, particularly prostate cancer. This gene and its protein can be used in peptide and gene therapy for cancer patients, as well as being useful as diagnostic tools (both for cancer sufferers and those with a predisposition to the disease) and in the production of cancer drugs
                                                                                                                                                                                                                                                                                    Ribozyme; hairpin; hammerhead; gene therapy; vasotropic; restenosis; ss.
                                                                                                               Gaps
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                                                                                         Query Match
3.0%; Score 12.6; DB 1; Length 19;
Best Local Similarity 78.9%; Pred. No. 4.4e+02;
Matches 15; Conservative 0; Mismatches 4; Indels
                                                                     G; 0 T; 0 U; 0 Other;
                                                                                                                                              CCACTCAGAGGAGTCTCTG 68
                                                                                                                                                                                                                                                                    cdk4 ribozyme binding site #10.
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                                                                                                                                                                                                                                                                                                                                                                        99WO-US028772.
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                                                                          Sequence 19 BP; 7 A; 7 C; 5
                                                                                                                                                                                                        AAA82829 standard; DNA; 19
                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                               (IMMI) IMMISOL INC.
                                                                                                                                                                                                                                                                                                                               WO200032765-A2.
                                                                                                                                                                                                                                                                                                                                                                        06-DEC-1999;
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                                                                                                                                                                                                                                                                                                             Mammalia.
                                                                                                                                                                                                                           AAA82829;
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                                                                                                                                                                                    RESULT 522
AAA82829/c
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ô 'n The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CDKI, PCNA and Cyclin BI. Representative examples of ribozyme recognition sites are given in AAA82415 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is efficient in Gaps ö Query Match 3.0%; Score 12.6; DB 1; Length 19; Best Local Similarity 78.9%; Pred. No. 4.4e+02; Matches 15; Conservative 0; Mismatches 4; Indels Sequence 19 BP; 1 A; 7 C; 5 G; 6 T; 0 U; 0 Other; 75 GAGGGCGCGCAGTGGACA 93 19 GAGGCCACAAAGTGGCCA 1 restenosis treatment ઠ

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The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinases coher than cell-cycle dependent kinases CNKI, PKNA and CYCLIN B1.
Representative examples of ribozyme recognition sites are given in AAA82415 to AAA86787. The ribozyme recognition sites are given in hibbiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is efficient in restenosis treatment
                                                                                                                     Ribozyme; hairpin; hammerhead; gene therapy; vasotropic; restenosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                              New hairpin and hammerhead ribozyme for inhibiting restenosis, cleaves RNA encoding a cyclin or cell-cycle dependent kinase other than CDK1, PCNA and Cyclin B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Angiotensin-converting enzyme gene; ACE; polymorphism; polymorphic marter, cardiovascular diseases, mycoardial infarction; unstable angina; hypertension; atherosclerosis; stroke; prognosis; drug screening; treatment outcome; human; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human angiotensin-converting enzyme (ACE) PCR primer, SEQ ID NO:2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      / Match 3.0%; Score 12.6; DB 1; Length 19; Local Similarity 78.9%; Pred. No. 4.4e+02; les 15; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19 BP; 3 A; 8 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                     Tritz R, Welch PJ, Barber JR, Robbins JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 85; 109pp; English.
                                                                                      Cyclin F ribozyme binding site #221.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       321 gracradecadecadec 339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GIGCIGACGGAGGAGTACC 1
                                                                                                                                                                                                                                             99WO-US028772.
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AAA38202 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-AUG-2000 (first entry)
AAA84953 standard; DNA; 19
                                                        (first entry)
                                                                                                                                                                                                                                                                                                       (IMMU-) IMMUSOL INC.
                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-412314/35.
                                                                                                                                                                                  WO200032765-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                        04-DEC-2000
                                                                                                                                                                                                                                             06-DEC-1999;
                                                                                                                                                                                                                                                                            04-DEC-1998;
                                                                                                                                                                                                              08-JUN-2000.
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                             AAA84953;
                                                                                                                                                   Mammalia.
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Assessing cardiovascular status in humans involves comparing test polymorphic pattern comprising polymorphic positions within genes encoding specific proteins, with reference polymorphic pattern.

Example 1; Page 48; 126pp; English

Norberg LT, Andersson MK, Lindstrom PHR,

WPI; 2000-318010/27

99WO-IB001678 98US-0104286P.

AAA84953/c

14-OCT-1998; 14-OCT-1998; 13-OCT-1999;

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The invention relates to a novel method of assessing the cardiovascular status in an individual and to newly identified polymorphisms in the gratus in an individual and to newly identified polymorphisms in the gense encoding angiocensin-converting enzyme (ACB), angiotensing in the competion of angiotensing and type 1 (ATI) and type 2 (ATI), angiotensingen (ACI), renin, a didosterone synthase, endochbelin receptor type A and Deta-adenseryic creeptor type 1 (ATI) and type 2 (ATI), angiotensing the acceptor type A and 2. The method comprises determining the sequence at one or more polymorphic positions within these genes, and comparing the creeptors 1 and 2. The method comprises determining the sequence at one or more polymorphic position a population of individual sexhibiting a patent of determining the predisposition of an individual to cardiovascular disorders such as myocaxdial infarction, unstable angina, or useful for determining the predisposition of an individual to cardiovascular seatus of a patient given a predicting the likely cardiovascular status of anticivascular drugs creatment regimen comprising administration of cardiovascular drugs (e.g., ACE inhibitors) bate-adrenergic receptor antagonists (between the cardiovascular drugs and probes for detecting genetic polymorphisms or in molecular creaments of the genes comprising a polymorphic site may be used as provides a basis for predicting the outcome of atreatment regimen.

Creatments of the genes comprising a polymorphic site may be used as primary arrays for high inhoughput soreaning. The genes, and the proteins creament of they encode are useful in the acreeing of potential cardiovascular eliminates trial and error in selecting a treatment for a particular creament expense of an individual searching of potential can be evaluated to individual cardiovascular patient. It also provides the ablity to a treatment regimen Adverse results in a detecting a treatment group per each and every and adverse results of any entering the number of patientes requir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 19 BP; 7 A; 3 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          360 GACTTCCTCACTTTCCTGG 378
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19 GATTTCTTCACCTCCTGG 1
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Best Local Similarity 78.9
Matches 15; Conservative
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ID AAA0
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AC AAA0
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20-APR-2000,

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Novel phosphodiesterase and its gene for research on complex mechanism of intracellular information transfer.
                                                                                                                                                                                                                                                                                                                Sequences AAA09589-A08592 encode human phosphodiesterase 10 (PDE10) proteins AAB26853-B26856. Phosphodiesterase 10 and its gene are useful for research on the complex mechanism of intracellular information transfer. The invention includes a recombinant vector containing a PDE10 gene, and a cell transfermed with the vector. Sequences AAA09593-A09606 represent PCR primers used in the isolation of the PDE10 polynucleotide sequences of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human ACE, AGT and AT1 genes polymorphisms PCR primer SBQ ID NO: 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; genetic polymorphism; disease diagnosis; treatment; cancer; cardiovascular system; nervous system; glaucoma; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                            3.0%; Score 12.6; DB 1; Length 19; 78.9%; Pred. No. 4.4e+02; tive 0; Mismatches 4; Indels
                                                Phosphodiesterase 10; PDE10; human; PCR primer; ss
                        PCR primer SEQ ID 17 used in PDE10 identification.
                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 19 BP; 2 A; 7 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lindstrom PHR, Norberg LT, Jonsson L,
                                                                                                                                                                                                                                                                                             Example 6; Page 27; 29pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GGTGCACCTGGAGCAGGGC 281
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GGTCCACCTGGAAGAGCGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99US-0126046P.
99WO-IB000497.
99US-0126243P.
99US-00471890.
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                                                                                                                                                                                                      (TANA ) TANABE SEIYAKU CO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 78.9
                                                                                                                                                                                                                             WPI; 2000-605129/58.
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                                                                                                    JP2000224992-A.
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24-MAR-1999;
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                                                                                                                                                                               30-NOV-1998;
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                                                                                                                              15-AUG-2000.
                                                                            Synthetic.
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                                                                                                                                                            The present invention is related to methods for determining the polymorphic pattern of an individual and using the results to determine their risk of a number of diseases, including cancer, cardiovascular diseases, glaucoma and nervous system disorders such as depression and neurodegenerative diseases. In addition, the methods can be used to adetermine the effects of different types of treatment for individuals, and thus enables appropriate therapies to be prescribed. THe PCR primers shown in sequences AAC61201-C61371 were all used to demonstrate the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Single nucleotide polymorphism; SNP; human; genetic disease;
disease susceptibility; cardiovascular system; endocrine system;
neurological system; forensic testing; paternity testing; PCR primer; ss.
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of an
                                      Assessing disease status in individual by determining sequence(s) at one or more polymorphic positions within the human genes encoding the protein(s) involved in physiological pathway associated with treatment
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Best Local Similarity 78.9%; Pred. No. 4.4e+02;
Matches 15; Conservative 0; Mismatches 4; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Single nucleotide polymorphism PCR primer #688.
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                                                                                                                                Example 1; Page 55; 141pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                            360 GACTICCICACITICCIGG 378
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Lipshutz RJ, Patil N,
                                                                                                                                                                                                                                                                                                         methods of the invention
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Olaisson E,

WPI; 2000-638268/61.

135

117 AGCAAGTACGGCATGCTGG 1 AGCACGTGAGGCATTCTGG

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AAC71168 standard; DNA; 19

AAC71168

AAC71168;

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The present invention is concerned with a number of human single nucleotide polymorphisms (SNPs) which the inventors identified in human genes. These SNPs can be used in disease diagnosis and prediction of an individual's susceptibility to disease, in forensic and paternity testing and in genetic mapping. In particular, the SNPs of the invention can be used to disgance susceptibility to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, echizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Single nucleotide polymorphism; SNP; human; genetic disease; disease susceptibility; cardiovascular system; endocrine system; neurological system; forensic testing; paternity testing; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleic acid selected from one of 106 genes comprising single nucleotide polymorphisms, allele-specific oligonucleotides to the genes are useful for phenotypic correlations, forensics, paternity testing, medicine and genetic analysis.
individual's susceptibility to disease, in forensic and paternity testin and in genetic mapping. In particular, the SNBs of the invention can be used to disgnose susceptibility to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, schizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's
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Lipshutz RJ, Patil N, Sklar P;
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                                                                                                                                                                     Query Match 3.0%; Score 12.6; DB 1; Length 19; Best Local Similarity 78.9%; Pred. No. 4.4e+02; Matches 15; Conservative 0; Mismatches 4; Indels
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                                                                                                                                     Seguence 19 BP; 4 A; 4 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Single nucleotide polymorphism PCR primer #720.
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Matches 15; Conservat
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                                                                                                                       Single nucleotide polymorphism; SNP; human; genetic disease; disease susceptibility; cardiovascular system; endocrine system; neurological system; forensic testing; paternity testing; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleic acid selected from one of 106 genes comprising single nucleotide polymorphisms, allele-specific oligonucleotides to the genes are useful for phenotypic correlations, forensics, paternity testing, medicine and
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Lipshutz RJ, Patil N, Sklar P;
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                                                                                             Single nucleotide polymorphism PCR primer #666.
                                                                                                                                                                                                                                                                                                                                                           (WHED ) WHITEHEAD INST BIOMEDICAL RES (APPY-) APFYMETRIX INC.
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Matches
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ID AAC7
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AC AAC7
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The present invention is concerned with a number of human single mucleotide polymorphisms (SNPs) which the inventors identified in human genes. These SNPs can be used in disease diagnosis and prediction of an individual's susceptibility to disease, in forensic and paternity testing and in genetic mapping. In particular, the SNPs of the invention can be used to diagnose susceptibility to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, echizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's
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                                Single nucleotide polymorphism; SNP; human; genetic disease;
disease susceptibility; cardiovascular system; endocrine system;
neurological system; forensic testing; paternity testing; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleic acid selected from one of 106 genes comprising single nucleotide polymorphisms, allele-specific oligonucleotides to the genes are useful for phenotypic correlations, forensics, paternity testing, medicine and genetic analysis.
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Single nucleotide polymorphism PCR primer #724.
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Lipshutz RJ, Patil N, Sklar P;
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Best Local Similarity 78.9
Matches 15; Conservative
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The present invention is concerned with a number of human single incleotide polymorphisms (SNPs) which the inventors identified in human genes. These SNPs can be used in disease diagnosis and prediction of an individual's susceptibility to disease, in forensic and paternity testing and in genetic mapping. In particular, the SNPs of the invention can be used to diagnose susceptibility to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, schizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's
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Sklar P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3.0%; Score 12.6; DB 1; Length 19; 78.9%; Pred. No. 4.4e+02; tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19 BP; 4 A; 4 C; 7 G; 4 T; 0 U; 0 Other;
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                                                                                           (WHED ) WHITEHEAD INST BIOMEDICAL RES (AFFY-) AFFYMETRIX INC.
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Patil N, Sl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-611722/58.
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Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               genetic analysis.
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                                                                                                                                                                                               Altshuler D,
Lipshutz RJ,
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Lipshutz RJ,
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Sequence 19 BP; 4 A; 4 C; 7 G; 4 T; 0 U; 0 Other;

diseases

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The present invention is concerned with a number of human single nucleotide polymorphisms (SNPs) which the inventors identified in human genes. These SNPs can be used in disease diagnosis and prediction of an individual's susceptibility to disease, in forensic and paternity testing and in genetic mapping. In particular, the SNPs of the invention can be used to diagnose susceptibility to diseases of the ardiovascular, endocrine and neurological systems, such as coronary artery disease, echizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's
for phenotypic correlations, forensics, paternity testing, medicine and
genetic analysis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3.0%; Score 12.6; DB 1; Length 19; 78.9%; Pred. No. 4.4e+02; tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19 BP; 4 A; 4 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        117 AGCAAGTACGGCATGCTGG 135
                                                                                                                Claim 8; Fig 5; 214pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
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Gaps

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Single nucleotide polymorphism; SNP; human; genetic disease;
disease susceptibility; cardiovascular system; endocrine system;
neurological system; forensic testing; paternity testing; PCR primer; ss.
                                                                       Single nuclectide polymorphism PCR primer #700.
AGCACGTGAGGCATTCTGG 19
                                  BB
                                  AAC71219 Standard; DNA; 19
                                                           (first entry)
                                                                                                               Homo sapiens.
                                                           09-FEB-2001
                                               AAC71219;
                      RESULT 533
                             AAC71219
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Altshuler D, Cargill M, Daley GQ, Ireland JS, Lander ES; Lipshutz RJ, Patil N, Sklar P; (WHED ) WHITEHEAD INST BIOMEDICAL RES (AFFY-) AFFYMETRIX INC.

30-MAR-2000; 2000WO-US008440.

WC200058519-A2.

05-CT-2000

99US-0127248P

31-MAR-1999;

WPI; 2000-611722/58.

Nucleic acid selected from one of 106 genes comprising single nucleotide polymorphisms, allele-specific oligonucleotides to the genes are useful for phenotypic correlations, forensics, paternity testing, medicine and genetic analysis.

Claim 8; Fig 5; 214pp; English.

The present invention is concerned with a number of human single nucleotide polymorphisms (SNPs) which the inventors identified in human genes. These SNPs can be used in disease diagnosis and prediction of an individual's susceptibility to disease, in forensic and paternity testing and in genetic mapping. In particular, the SNPs of the invention can be used to diagnose susceptibility to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, schizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention is concerned with a number of human single nucleotide polymorphisms (SNPs) which the inventors identified in human genes. These SNPs can be used in diesase diagnosis and prediction of an individual's susceptibility to disease, in forensic and paternity testing and in genetic mapping. In particular, the SNPs of the invention can be used to diagnose ausceptibility to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, schizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's
                                                                                                                                                                                                                                                                                                                     Single nucleotide polymorphism; SNP; human; genetic disease; disease susceptibility; cardiovascular system; endocrine system; neurological system; forensic testing; paternity testing; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nucleic acid selected from one of 106 genes comprising single nucleotide polymorphisms, allele-specific oligonucleotides to the genes are useful for phenotypic correlations, forensics, paternity testing, medicine and
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Score 12.6; DB 1; Length 19;
Pred. No. 4.4e+02;
0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 19 BP; 4 A; 4 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                         Single nucleotide polymorphism PCR primer #686.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     117 AGCAAGTACGGCATGCTGG 135
                                                                        117 AGCAAGTACGCCATGCTGG 135
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                                                                                                       1 AGCACGTGAGGCATTCTGG 19
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     3.0%;
Best Local Similarity 78.9%;
Matches 15; Conservative
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                                                                                                                                                                                                                                                            (first entry)
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Lipshutz RJ,
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                                                                                                                                                                                                                          AAC71198;
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AAA65792 standard; DNA; 19 BP. 535 RESULT 53

AAA65792;

(first entry) 22-NOV-2000 Human leukocyte antigen C exon 3 sequencing primer SEQ ID NO:13

Human; WHL gene; sequencing; mutation; human leukocyte antigen; HLA; transplantation surgery; detection; identification; primer; pathogenic microorganism; ss.

Homo sapiens

IS6083699-A.

04-JUL-2000

98US-00009483 10-JAN-1998;

96US-00640672. 96US-00684498. 97US-00807138. 97WO-US007134. 01-MAY-1996; 19-JUL-1996; 27-FEB-1997; 29-APR-1997;

(VISI-) VISIBLE GENETICS INC.

Dunn JM, Larson MT, Lacroix J, Shipman R, Leushner J; Hui M,

WPI; 2000-464336/40.

Bi-directional sequencing of nucleic acid polymers for identifying pathogens or detecting mutations by using a single reaction mixture having first and second primers with different, spectroscopically-distinguishable labels.

Example 2; Col 11; 27pp; English.

The present invention describes a method for simultaneously determining the position of a nucleotide base in a target region of both strands of a denatured duplex nucleic acid polymer. The method comprises using a single set of reaction mixture that is combined with the nucleic acid polymer. The reaction mixture contains first and second oligonucleotide polymer. The reaction mixture contains first and second oligonucleotide primers, each with different, spectroscopically-distinguishable contained in the method is used to detect mutations, especially medically significant mutations, in samples derived from a human patient, animal, plant or microorganism, and for the determination of human contains first mutations, and for the determination of human also be used to detect and identify microorganisms, especially pathogenic microorganisms, in a sample, and in in situ sequencing pathogenic microorganisms, in a sample, and in in situ sequencing can then removed from a selected location on the tissue preparation and loaded onto a gel for sequence analysis. The sequencing reaction is contone of a disease condition is known, but the causative mutation is cont. The present sequence a sequencing primer for the human cont. The present sequence as sequencing primer for the human contoner. The present invention is manyle from the present invention 

Seguence 19 BP; 2 A; 6 C; 11 G; 0 T; 0 U; 0 Other;

0; Gaps 3.0%; Score 12.6; DB 1; Length 19; 78.9%; Pred. No. 4.4e+02; tive 0; Mismatches 4; Indels 3.0 Best Local Similarity 78.9 Matches 15, Conservative

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RESULT 536 AAC85428

AAC85428 standard; cDNA; 19 BP. 

AAC85428;

(first entry) 20-APR-2001 Primer oML69 amplifies Salmon MHC class II beta and alpha promoters

Promoter; regulation; expression; MHC class II; RAN; U2A'; primer; immune response; productivity; DNA vaccination; cytokine; amplify; interferon gamma; beta-carotene; polymerase chain reaction; PCR; ss

Synthetic.

WO200077232-A1.

21-DEC-2000.

09-JUN-2000; 2000WO-NO000202.

10-JUN-1999; 99NO-00002819

(GENO-) GENOMAR AS.

Syed M, Lundin M;

WPI; 2001-080695/09.

Novel promoters from Atlantic salmon for regulating expression of nucleotide constructs, as DNA vaccines for protecting salmon and other fish species against viral, bacterial infections.

Example 1; Page 7; 29pp; English.

The sequences given in AAC85425-28 are primers which were used in the amplification of the Salmon MHC class II bees and alpha promoters and introduced in recorded in nucleotide constructs for the purpose of regulating the expression of such constructs for the purpose of regulating the expression of such constructs. For example, the salmon MHC class II beta-promoter was constructs. For example, the salmon MHC class II beta-promoter was inserted in a plasmid vector carrying the lacz gene. The resulting plasmid vactor to promote transcription of the lacz gene. The resulting plasmid vactor different higher than the mean value +2 containing plasmid, showed an ELISA titer higher than the mean value +2 containing plasmid, showed an ELISA titer higher than the mean value +2 containing plasmid, showed an ELISA titer higher than the mean value +2 containing plasmid, showed an ELISA titer higher than the mean value +2 containing plasmid, showed an ELISA titer higher than the mean value +2 containing plasmid, showed an ELISA titer higher than the mean value +2 constructs such as these, may be used in vivo to achieve productivity constructs such as these, may be used in vivo to achieve productivity constructs such as these, may be used in vivo to achieve productivity constructs may also be useful for DNA vaccination of salmon and other constructs may also be useful for DNA vaccination of salmon and other constructs may also be useful for DNA vaccination of salmon and other in somatic tissues and may also be combined with MHC alleles for optimizing the expression and presentation of pathogenic antigens. Promoters may also be used to influence gene expression, for regulating cuttives and seroage of beta-canotenes and compounds responsible for meat cuttive and scrage of beta-canotenes and compounds responsible for meat coloring in salmonids. The promoters is active. The constitutive coloring the expression the promoter is active. The constitutive constitutive in allenged to influence promoter is active. The constitutive consti

Sequence 19 BP; 4 A; 4 C; 6 G; 5 T; 0 U; 0 Other;

ö Query Match
3.0%; Score 12.6; DB 1; Length 19;
Best Local Similarity 78.9%; Pred. No. 4.4e+02;
Matches 15; Conservative 0; Mismatches 4; Indels

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Gaps

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The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a ribozyme (1) which cleaves RNA encoding a cytckine involved in inflammation, matrix metalloproteinase (MMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid segment encoding (1). (1) can have antipsoriatic, dermatological, cytostatic, antiseborrheic, antidiabetic, antisickling, cleaves RNA encoding (1). (1) can have antipsoriatic, and segment encoding (1). (2) can have antipsoriatic, cleaves RNA encoding cytokine involved in inflammation. (1) can be used in gene therapy. (1) and (II) are useful for treating proliferative skin diseases such as psoriasis, atopic dermatitis, actinic keratosis, equamous or basal cell carinoma and viral or sebornheic wart. They can also be used for treating proliferative eye diseases such as diabetic retinopathy, virreoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal detechment, and for treating and preventing prematurity and retinal detechment, and for treating and preventing scar. AAH57877 to AAH62099 sepresent sequences used in the scar. AAH57877 to AAH62099 sepresent sequences used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent kinases.
cytokine, inflammation, cell-cycle dependent kinase, cyclin; MMP; matrix metalloproteinase; growth factor; reductase, scarring, cytostatic, antipsoriatic, dermatological, antiseborrheic; antidabetic; virucide; antifickling; ophthalmological; keratolytic; gene therapy; viral wart; appid dermatitis, actinic keratosis; squamous cell carcinoma; basal cell carcinoma; seborrheic wart; vitreoretinopathy; scar; sickle cell retinopathy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                / Match
3.0%; Score 12.6; DB 1; Length 19;
Local Similarity 78.9%; Pred. No. 4.4e+02;
les 15; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 19 BP; 1 A; 7 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 102; 408pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             75 GAGGGCGCGCACAGTGGACA 93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            н
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tritz R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (IMMU-) IMMUSOL INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-300427/31.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Robbins JM,
                                                                                                                                                                                                      sapiens
                                                                                                                                                                                                                                                                                                                             03-MAY-2001:
                                                                                                                                                                                                                          Synthetic.
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence is a PCR primer for wild-type leuk gene from E.coli, which encodes alpha-isopropylmalate synthase (IPMS). The leuk gene was used to generate a mutant alpha-IPMS, which is de-sensitised in feedback inhibition by L-leucine. The mutant alpha-IPMS is useful for the production of L-leucine, which is useful for medical treatment, as a pharmaceutical or in the chemical industry or as a growth factor useful for production of other amino acids such as 19sine. The present sequence was used to amplify the leuk gene for use in the present invention
                                                                                                                                                                                                                                                                                                                                                  Alpha-isopropylmalate synthase; enzyme; IPMS; leucine production; leuA; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cell-cycle dependent kinase cdk4 ribozyme binding site SEQ ID NO:415.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New polypeptide with alpha-isopropylmalate synthase activity and decreased feedback inhibition of activity by L-leucine, useful for production of L-leucine for medical treatment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3.0%; Score 12.6; DB 1; Length 19; 78.9%; Pred. No. 4.4e+02; ive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lunts MG, Kozlov YI, Ivanovskaya LV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19 BP; 2 A; 4 C; 8 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 1; Page 9; 19pp; English
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                                                                                                                                                       AAF31028 standard; DNA; 19 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAH57991 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                      leuA gene PCR primer LeuA9
                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ouery Match
Best Local Similarity 78.9 Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (AJIN ) AJINOMOTO KK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-125730/14.
                                                                                                                                                                                                                                                                                                                                                                                                                           Escherichia coli.
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Gusyatiner MM, Voroshilova EB;

09-JUL-1999;

EP1067191-A2.

.0-JAN-2001.

05-APR-2001

AAF31028;

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Gaps

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Human; ribozyme therapy; hairpin ribozyme; hammerhead ribozyme;

Cyclin F ribozyme binding site SEQ ID NO:2539.

Human, ribozyme therapy, hairpin ribozyme; hammerhead ribozyme; recognition site, target, ribozyme binding site, eye disease; vulnerary; proliferative disease; skin disease; psoriasis; diabetic retinopathy;

10-SEP-2001

RESULT 538
AAH57991/C
ID AAH57991/C
XX
AC AAH5.
DT 10-S1
XX
XX
XX
Human
KW FOOO

AAH57991;

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538

Homo sapiens. Synthetic.

26-OCT-1999;

03-MAY-2001

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The present invention relates to a number of prostate-specific sequences derived from the human PS118 gene. These can be used in the detection, monitoring and treatment of prostate diseases, particularly prostate cancer. The PS118 fragments of the invention were isolated from a prostate tissue expressed sequence tag (EST) library. The present sequence is a PCR primer used to isolate a sequence of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel PS118 polypeptide for detecting, diagnosing, staging, monitoring, prognosticating, preventing, treating, or determining predisposition of individual to diseases and conditions of prostate, e.g. prostrate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human, mouse; HPC2; prostate cancer; neoplastic growth; cytostatic; si
gene therapy; prostate cancer predisposing gene; chimpanzee; gorilla;
sequencing primer; PCR primer.
                       Human, prostate, prostate-specific sequence, prostate cancer, PS118, cytostatic, gene therapy, PCR, primer, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human prostate cancer predisposing gene (HPC2) PCR primer #95.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 3.0%; Score 12.6; DB 1; Length 19; Best Local Similarity 78.9%; Pred. No. 4.4e+02; Matches 15; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Colpitts TL, Friedman PN,
Klass MR, Kratochvil JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 19 BP; 3 A; 2 C; 10 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 2; Page 41; 58pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        139 GCCTGGCGGTGGAGGCCGG 157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 GACTGGCGGTAGAGGTTGG 19
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                                                                                                                                                                                                                                                    26-NOV-2001; 2001US-00991681.
                                                                                                                                                                                                                                                                                                    97US-00842385.
98US-00065383.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Billingel PA, Cohen M,
Granados EN, Hodges SC,
Russell JC, Stroupe SD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (KLAS/) KLASS M R.
(KRAT/) KRATOCHVIL J D
(ROBE) ROBERLS - RAPP L
(RUSS/) RUSSELL J C
(STRO/) STROUPE S D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GORDAN J.
GRANADOS E N.
HODGES S C.
                                                                                                                                                                                                                                                                                                                                                                               BILLINGEL P A.
                                                                                                                                                                                                                                                                                                                                                                                                                            COLPITIS T L.
FRIEDMAN P N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-665429/71.
                                                                                                                                                       US2002086316-A1.
                                                                                                                                                                                                                                                                                                                                                                                                          COHEN M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Granados EN,
Russell JC,
                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                       23-APR-1997;
23-APR-1998;
                                                                                                                                                                                                       04-JUL-2002
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                                                                                                                                                                                                                                                                                                                                                                            (BILL/)
                                                                                                                                                                                                                                                                                                                                                                                                                                  (COLP/)
(FRIE/)
(GORD/)
(GRAN/)
(HODG/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 541
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HOMO
     8XXXXXXXXXXXX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ઠ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a ribozyme (I) which cleaves RNA encoding a cytokine involved in tillammation, matrix metalloproteinase (WMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid molecule, antisoching, antidiabetic, antisickling, ophthalmological, cytostatic, antiseborrheic, antidiabetic, antisickling, dermatological, vulnerary, keratolytic and virucide activities, and cleaves RNA encoding cytokline involved in inflammation. (I) can be used in gene therapy. (I) and (II) are useful for treating proliferative skin diseases such as psoriasis, atopic dermatitis, actinic keratosis, squamous or basal cell carionoma and viral or seborific wart. They can also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal detachment, and for treating and preventing scarring such as keloid, adhesion and hypertrophic or hypertrophic burn scar. AMF3577 to AMH62099 represent sequences used in the exemplification of the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent kinases.
recognition site; target; ribozyme binding site; eye disease; vulnerary; proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin; MMP; matrix metalloproteinase; growth factor; reductase; scarring; cytostatic; antipsoriatic; dermatological; antiseborrheic; antidiabetic; virucide; antisickling; ophthalmological; keratolytic; gene therapy; viral wart; atopic dermatitis; actinic keratosis; squamous cell carcinoma; basal cell carcinoma; seborrheic wart; vitreoretinopathy; scar;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
3.0%; Score 12.6; DB 1; Length 19;
Best Local Similarity 78.9%; Pred. No. 4.4e+02;
Matches 15; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Seguence 19 BP; 3 A; 8 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 1; Page 256; 408pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             321 GIGCIGGCGGCGGACGACC 339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19 Grecreaceadadarace 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                       26-OCT-2000; 2000WO-US029500.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Robbins JM, Tritz R;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (IMMU-) IMMUSOL INC.
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Gordan J; Roberts-Rapp L;

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Gaps

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AAL49572 RESULT 540
AAL49572
ID AAL4957
AC AAL4957
XX
XX
DT 27-NOVDT 27-NOVDE Human g

ઠે g WPI; 2002-643344/69

Rommens JM;

(FARB ) BAYER AG

Smolyar A;

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G-protein coupled receptor; receptor; serotonin; 5-hydroxytryptamine; human; antibacterial; virucide; fundicide; protezoacide; neuroprotective; cardiant; antidepressant; hypertensive; hypotensive; diuretic; costeopathic; antiulcer; antiliamatory; antiallergic; cytostatic; nootropic; analgesic; gene therapy; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a human prostate cancer predisposing gene coding for an HPC2 polypeptide. The DNA and protein sequences are useful as diagnostic reagents for identifying a mutant HPC2 nucleotide sequence in a suspected mutant HPC2 allele by comparing the sequence of the suspected mutant HPC2 allele with a wild-type HPC2 sequence. The sequences are also useful for detecting an alteration in HPC2, where the alteration is associated with cancer in a human. The method involves analysing an HPC2 gene or an HPC2 gene expression product from a tissue of the human. The HPC2 gene is useful as a marker for prostate cancer and can be used in gene therapy techniques to suppress neoplastic growth of recipient cells which carry the mutant HPC2 and encoding human and mouse HPC2 and in the methods of the invention, CDNA encoding human and mouse HPC2 and
                                                                                                                                                                                                                                                                                         Novel nucleic acid sequence encoding HPC2 polypeptide, which is marker for prostate cancer, is useful in gene therapy techniques to restore HPC2 normal levels by which neoplastic growth is suppressed in recipient cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human serotonin-like G-protein coupled receptor PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19 BP; 7 A; 7 C; 5 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 in the methods of the invention, cDNA encoding cDNA encoding HPC2 paralogues and orthologues
                                                                                                                                                                                                                      Tavtigian SV, Teng DHF, Simard J,
                                                                                                                                                                                                                                                                                                                                                                         Example 8; Page 75; 239pp; English.
                                                                                                                                                              (MYRI-) MYRIAD GENETICS INC.
(HOSP-) HOSPITAL FOR SICK CHILDREN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CCACTCAGAGGAGTCTCTG 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ccacacadaddaddccacag 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-JAN-2001; 2001US-0264071P.
24-SEP-2001; 2001US-0324054P.
                                                                                         07-MAY-2001; 2001WO-US014602.
                                                                                                                           35-MAY-2000; 2000US-00564805.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity 78.9
nes 15; Conservative
                                                                                                                                                                                                                                                          WPI; 2002-066599/09.
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                   WO200185911-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-NOV-2002
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                                                     15-NOV-2001
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                                                                                                                                                                                                                               The present sequence is a forward primer for a novel human serotonin-like G-protein coupled receptor (5HT-like GPCR, see ABN64695). The primer was used in the RT-PCR amplification of 5HT-like GPCR mRNA in order to determine the expression profile of the receptor. 5HT-like GPCR mRNA was highly expressed in cerebellum, postcentral gyrus, dorsal root ganglia, erythrocytes, lung chronic obstructive pulmonary disease (GOPD), oesophagus, ileum chronic inflammation, benign prostatic hypertrophy (BPH) and penis. The invention provides reagents which regulate the 5HT-like GPCR and reagents which bind to 5HT-like GPCR gene products. These reagents can play a role in preventing, ameliorating or correcting dysfunctions or diseases including GOPD, a cardiovascular disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                         cancer, a urinary disorder, obssity, diabetes, a central nervous system (CNS) disorder, asthma or a haematological disorder (all claimed) in a subject. The reagent is especially an antisense oligonucleotide, ribozyme or antibody. Pharmaceutical compositions comprising the reagent, or an expression vector encoding 5HT-like GPCR, are claimed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome;
                                                                                                          New G-protein coupled receptor (GPCR) polynucleotide and its encoded procein, useful for identifying modulators of GPCR activity, and in gene therapy for treating bacterial infection, cancer, acute heart failure or Parkinson's disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3.0%; Score 12.6; DB 1; Length 19; ilarity 78.9%; Pred. No. 4.4e+02; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human chromosome 1p36-35 PCR primer SEQ ID NO:2078.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 19 BP; 2 A; 7 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                     Example 21; Page 124; 164pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           284 CACCAAGCTGGTGAAGGAC 302
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (RIKA ) RIKAGAKU KENKYUSHO.
(GENO-) GENOTEX YG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABL45034 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Arraying genome clones.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-144136/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 JP2001321190-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15;
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Matches
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Gaps

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3.0%; Score 12.6; DB 1; Length 19; 78.9%; Pred. No. 4.4e+02; tive 0; Mismatches 4; Indels

WPI; 2002-187683/24.

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The present invention describes a method of arraying genome clones. The method comprises: (a) clones of the genomic libraries contained in method comprises: (a) clones of the genomic libraries contained in multiwell plates numbered for discrimination are mitsed in each of the multiwell plates; (b) a primer designed based on the chromosome marker sequence is added to the mixture to carry out an amplification reaction; (c) a signal corresponding to the marker is detected from the resultant complified product to specify the discrimination Nos. of the multiwell plates containing the clones having said marker sequence; (d) the order of the markers is changed so that the same discrimination Nos. succeed to the maximum in the specified discrimination Nos. to array the multiwell plates; (e) the clones in the multiwell plates of the specified discrimination Nos. are mixed respectively in each wells of longitudinal and lateral directions; (f) the multiwell plates of the specified contains are cultured and the amplified products; (h) the clones in the multiwell plates are specified from the detected result; and (i) the clones are reconstituted as the positions on the chromosome and arrayed. The microarray is useful for gene analysis. ABM42957 to ABM43231 to ABM45634 creptesent PCR primers for human chromosome light., which are specifically claimed for use in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PS118; prostate; marker; prostate cancer; human; sequencing; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gordon J;
Roberts-Rapp L;
                                                                                                                                                                                                                                                                                                                                                                                                                               .
0
                                                                                                                                                                                                                                                                                                                                                                                            3.0%; Score 12.6; DB 1; Length 19;
llarity 78.9%; Pred. No. 4.4e+02;
Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Copipitts TL, Friedman PN,
Klass MR, Kratochvil JD,
                                                                                                                                                                                                                                                                                                                                                                 Sequence 19 BP; 5 A; 7 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Prostate-specific PS118 clone sequencing primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              87 GTGGACATCACCACGTCTG 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 Grecacarcaccaracere 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABA91662 standard; DNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                97US-00842385
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98US-00065383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hodges SC, Stroupe SD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GRANADOS E N.
HODGES S C.
KLASS M R.
KRATOCHVIL J D.
ROBERTS-RAPP L.
RUSSELL J C.
STROUPE S D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BILLINGEL P A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COPLPITTS T L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FRIEDMAN P N.
                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
hes 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US2001055758-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COHEN M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Billingel PA,
Granados EN, F
Russell JC, St
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GORDON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-DEC-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABA91662;
                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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(COHE/)
(COPL/)
(FRIE/)
(GORD/)
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(KLAS/)
(KRAT/)
(ROBE/)
(RUSS/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 544
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The present sequence is that of a sequencing primer designed from sequencing information of a prostate-specific PS118 consensus sequence tages ABA91561). It was used in the sequencing of PS118 expressed sequence tag-specific clones (see ABA91642-S0) transcribed from human prostate tissue. PS118 polypeptides (see AAMS0809-13), polynucleotides are useful for and treating, or determining the predisposition of an individual to, hyperplasia, prostatitis, prostatic intraepithelial neoplasia, prostate cancer, tumours and metastases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lolium perenne; perennial ryegrass; plant; cell wall; lignification; cellulase; enzyme; lignin biosynthesis; cellulose degradation; CCOAMT; caffeoyl-CoA 3-O-methyltransferase; cinnamyl alcohol dehydrogenase; CBD; caffeic acid O-methyltransferase; OMT; cinnamete-4-hydroxylase; CAH; cinnamoyl-CoA reductase; CCR; percxidase; PER; ferulate-5-hydroxylase; F5H; CELL; phenylalanine ammonia lyase; PAL; 4-coumarate:COA ligase; 4CL; ryegrass; fescue species; molecular genetic marker; PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel nucleic acid encoding lignification and cellulase enzymes or their related enzymes useful for modifying lignin biosynthesis and cellulose degradation in plants to manipulate plant cell wall.
                              Detecting presence of target PS118 polynucleotide in test sample, useful for detecting, diagnosing, staging, monitoring, prognasticating, preventing or treating or determining predisposition to prostate disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                                                                                                    3.0%; Score 12.6; DB 1; Length 19; 78.9%; Pred. No. 4.4e+02; tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ong EK, Emmerling M;
                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 19 BP; 3 A; 2 C; 10 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (AGRI-) AGRIC VICTORIA SERVICES PTY LTD. (AGRE-) AGRESEARCH LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lolium perenne LpPeroxidasel primer #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 3; Page 37; 436pp; English
                                                                                                                  Example 2; Page 41; 57pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          139 GCCTGGCGGTGGAGGCCGG 157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        | ||||||||| |||| ||| ||| GACTGGCGGTAGAGGTTGG 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Spangenberg G, Sawbridge TI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABN87259 standard; DNA; 19 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-SEP-2001; 2001WO-AU001221.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-SEP-2000; 2000AU-00000419.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 78.9
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-444025/47.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lolium perenne
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04-APR-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABN87259;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 545
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABN87259
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The present invention describes a nucleic acid (1) or its fragment encoding caffeoyl-CoA 3-O-methyltransferase (CCAMT), cinnamyl alcohol dehydrogenase (CAD), caffeid acid 0-methyltransferase (OMT), cinnamate-4-hydroxylase (CRE), peroxidase (ORE), orinnamate-4-hydroxylase (CRE), peroxidase (PER), cinnamate-4-lydroxylase (CRE), peroxidase (PER), cinnamate-4-lydroxylase (FSH), phenylalanine ammonia lyse (Dolium perenne) or fescue species (TI), its nucleotide sequence information and/or single nucleotide polymorphisms is useful as a molecular genetic marker. (I) its nucleotide sequence information and/or callulose degradation in a plant to manipulate cell biosynthesis and/or callulose degradation in a plant to manipulate cell walls. (I) or its fragments are useful for isolating cDNAs and genes conciding homologous proteins from the same or other plant species, as hybridisation probes to screen libraries from the desired plant. Short segments of (I) or its fragment are useful in amplification proteocls to amplify longer nucleic acids or its fragments encoding homologous genes from DNA or RNA. (I) or its fragments are useful as molecular genetic markers for quantitative trait loci (QTL) tagging, QTL mapping, DNA ingerprinting, and in marker assisted selection, particularly in ryegrass and fescues, and in forage and turf grass improvement, e.g. cagging QTLs for herbage quality traits, dry matter digestibility, mechanical stress tolerance, disease resistance, insect pest resistance, plant stature, leaf and stem colour. AbN87220 to AbN87272 represent prometrion

Sequence 19 BP; 8 A; 7 C; 3 G; 1 T; 0 U; 0 Other;

; 3.0%; Score 12.6; DB 1; Length 19; Similarity 78.9%; Pred. No. 4.4e+02; Conservative 0; Mismatches 4; Indels 392 CGCCAAGAAGGTCTTCTAC 410 1 CGCCAAGAAGAACCTCAAC 19 3.03 Best Local Similarity 78.9° Matches 15, Conservative 셤 ઠે

RESULT 546 ABZ76924

ABZ76924 standard; DNA; 19 BP 07-MAY-2003 (first entry) ABZ76924; 

Acyl CoA:diacylglycerol transferase; DGAT; enzyme; chromosome 8; human; milk; meat marbling; low fat; polymorphic; SNP; single nucleotide polymorphism; PCR primer; ss.

Homo sapiens. Synthetic.

WO2003004630-A2.

16-JAN-2003.

Fries H, Winter A;

New nucleic acid molecule comprising a sequence of an allele of a polymorphic bovine acyl CoA-diacylglycerol transferase gene useful for testing a mammal for its predisposition for fat content of milk and for testing a mamm meat marbling.

Example 2; Page 27; 91pp; English.

The present invention describes a nucleic acid molecule (NA) (I) encoding to indicative for low fat content of milk and to low meat marbling to or indicative for low fat content of milk and to low meat marbling or (intramuscular fat content). Human DGAT is located to chromosome 8, and (intramuscular fat content). Human DGAT is located to chromosome 8, and mammal for its predisposition for fat content of milk and/or its content of predisposition for meat marbling. The method comprises analysing the gene or content of milk and/or its spredisposition for meat marbling. The method comprises analysing the gene conding DGAT for nucleotide polymorphisms (e.g. single nucleotide polymorphisms are located in the coding region of the DGAT contention polymorphisms are located in the coding region of the DGAT contelled has at the position 10433 and 10434 of the DGAT contelled has at the position 10433 and 10434 of the DGAT contelled has at the position 10433 and 10434 of the DGAT content of milk and low meat marbling. The nucleic acid molecule has at the position for low fat content of milk and low meat marbling. The nucleic acid molecule has at the position for low fat content of corresponding to position 10433 and 10434 of the DGAT corresponding to position 10433 and 10434 of the DGAT gene high correlate with a predisposition for high content of milk and high meat marbling. The nucleic acid molecule has at the position of region which is responsible for the regulation of the expression of the present content of supposition which is responsible for the regulation of the expression of the present the position of the expression of the present for supposition the exemplification of the present invention

Sequence 19 BP; 2 A; 7 C; 5 G; 5 T; 0 U; 0 Other;

0; Gaps Query Match
3.0%; Score 12.6; DB 1; Length 19;
Best Local Similarity 78.9%; Pred. No. 4.4e+02;
Matches 15; Conservative 0; Mismatches 4; Indels

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Gaps

ACC62358 standard; DNA; 19 BP. RESULT 547 ACC62358 

ACC62358;

23-JUN-2003 (first entry)

Human NOV5 forward PCR primer SEQ ID NO:233.

Human; NOVX; antiatherosclerotic; hypotensive; cardiant; dermatological; anorectic; immunosuppressive; cytostatic; antidiabetic; antidiabetic; antidiabetic; antidiabetic; antidiabetic; antidiabetic; antidiabetic antidiabetic antidiabetic constructive; nootropic; antiparkinsonian; metabolic; antidipaemic; gene therapy; cardiomyopathy; atherosclerosis; hypertension; scleroderma; congenital heart defect; aortic stenosis; valve disease; transplantation; tuberous sclerosis; obesity; congenital adrenal hyperplasta; diabetes; prostate cancer; metabolic disorder; neoplasm; lymphoma; uterus cancer; fertility; haemophila; hypercoaqulation; graft versus host disease; idiopathic thrombocytopenic purpura; Albs; bronchial asthma; anorexia; diopathic thrombocytopenic purpura; Albs; bronchial asthma; anorexia; cancer-associated cacheria; Albrimer's disease; parkinson's disease; cancer-associated cacheria; Albrimer's disease; parkinson's disease; immune disorder; haematopoietic disorder; dyslipidaemia metabolic syndrome X; PCR primer; ss.

Homo sapiens.

Synthetic

WO2003023001-A2

20-MAR-2003.

09-SEP-2002; 2002WO-US028538.

07-SEP-2001; 2001US-0318120P.

rng.res

Human DGAT gene forward PCR primer 1534.

05-JUL-2002; 2002WO-EP007520.

06-JUL-2001; 2001EP-00116412. 13-MAY-2002; 2002US-0379412P.

(ARBE-) ARBEITSGEMEINSCHAFT DEUT RINDERZUECHTER.

WPI; 2003-239205/23.

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10-SEP-2001; 2001US-0318430P.
17-SEP-2001; 2001US-0322636P.
17-SEP-2001; 2001US-0322816P.
17-SEP-2001; 2001US-0322816P.
19-SEP-2001; 2001US-0322816P.
20-SEP-2001; 2001US-032361P.
20-SEP-2001; 2001US-032363P.
25-SEP-2001; 2001US-032363P.
25-SEP-2001; 2001US-032499P.
14-DEC-2001; 2001US-032499P.
14-DEC-2001; 2001US-032499P.
14-DEC-2001; 2001US-032699P.
16-DEC-2001; 2001US-032999P.
17-WAY-2002; 2002US-039999P.
17-WAY-2002; 2002US-038988P.
                                                                                                        2002US-0396412P
                                                                                                               13-AUG-2002; 2002US-0403517P.
06-SEP-2002; 2002US-00236417.
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Wed ADF 21 12:38:21 2004

### (CURA-) CURAGEN CORP.

Agee ML, Alsobrook JP, Anderson DW, Berghs C, Boldog FL;
Burgess CE, Casman SJ, Catterton B, Chant JS, Chaudhuri A;
Crabtree J, Dipippo VA, Edinger SR, Eisen AJ, Ellerman K,
Gangolli EA, Gerlach VL, Giot L, Gorman L, Guo X, Gusev VY, Ji W;
Kekuda R, Khramtsov NV, Leach MD, Lepley DM, Li L, Liu X;
Malyankar UM, Miller CE, Ooi CE, Ort T, Padigaru M, Patturajan M;
Pena CEA, Rieger DK, Rothenberg ME, Shenoy SG, Shimkets RA;
Spaderna SK, Spytek KA, Taupier RJ, Twomlow N, Vernet CAM, Voss EZ;
Zerhusen BD, Zhong M;

#### WPI; 2003-313241/30.

Novel human proteins and mucleic acid encoding the proteins, useful for diagnosis, treatment and prevention of disorders involving the human protein or nucleic acid e.g. cardiac and neurological disorders.

### Example C; Page 301; 460pp; English.

The present invention describes isolated human NOVX proteins, where X is

1 to 42. ACC62236 to ACC62345 encode the human NOVX proteins given in
ARS64167 to ABR54276. NOVX sequences have antiatheroscelectic, cardiant,
hypotensive, dermatological, anorectic, immunosuppressive, cytostatic,
antidiabetic, antiinfertility, haemostatic, antiinflammatory, anti-HIV,
antidiabetic, metabolic, immunomodulator, neuroprotective, nootropic,
antidabetic, metabolic, immunomodulator, neuroprotective, nootropic,
antiparkinsonian and antilipaemic activities, and can be used in gene
therapy. NOVX proteins are useful for treating or preventing a pathology
associated with the human disease. NOVX nucleic acids, proteins and
antibodies can be used in the treatment and diagnosis of cardiomyopathy,
atheroaclerosis, hypertension, congenital heart defects, acrid estnosis,
congenital adrenal hyperplasia, prostate cancer, diabetes, metabolic
disorders, neoplasm, lymphoma, uterus cancer, fertility, haemophilia,
hypercoagulation, idiopathic thrombotycopenic purpura, graft versus host
disease, AlDS, bronchial asthma, cromm's disease, multiple sclerosis,
infectious disease, parkinson's disease, multiple sclerosis,
confernieral adrenal hyperplasia, cancer-associated cachexia, cancer,
disease, AlDS, bronchial asthma, cromm's disease, multiple sclerosis,
hyperometer, which are used in examples from the present invention.
ARS54277 represents a human trvaningen protein diven in commarison with ABK54277 represents a human trypsinogen protein given in comparison with the human NOV35b protein in the exemplification of the present invention

# Sequence 19 BP; 6 A; 3 C; 9 G; 1 T; 0 U; 0 Other;

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Gaps
                                                               ö
Query Match
3.0%; Score 12.6; DB 1; Length 19;
Best Local Similarity 78.9%; Pred. No. 4.4e+02;
Matches 15; Conservative 0; Mismatches 4; Indels
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short interfering nucleic acid; siNA; downregulation; inhibition; mitogen-activated protein kinase; MAP kinase; MAPK; RNA interference; cytostatic; ancrediabetic; antiinflammatory; antiasthmatic; immunosuppressive; antibacterial; antiinflammator; antiarthritic; antipacriatic; gasrointestinal; obesity; diabetes; tumour; inflammatory disease; asthma; septic shock; rheumatoid arthritis; psoriasis; inflammatory bowel disease; drug screening; genetic engineering; pharmacogenomic; gene mapping; ss.
                                                                                                                                 Mitogen activated protein kinase sina oligonucleotide SEQ ID NO:414
260 caccercacercacae
         1 CAGGGAGGACCTGGAGAAG 19
                                                                                                                                                                                                                                                                                                                                         11-MAR-2002; 2002US-0363124P.
06-UN-2002; 2002US-036782P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-040293P.
15-JAN-2003; 2003US-0440129P.
                                                                     ADE29792 standard; RNA; 19 BP.
                                                                                                                                                                                                                                                                                                                               20-FEB-2002; 2002US-0358580P.
                                                                                                                                                                                                                                                                                                           28-JAN-2003; 2003WO-US002510
                                                                                                             29-JAN-2004 (first entry)
                                                                                                                                                                                                                                                                    WO2003072590-A1.
                                                                                                                                                                                                                                                                                        04-SEP-2003.
                                                                                                                                                                                                                                                Synthetic.
                                                                                         ADE29792;
                                                                               용
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New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of mitogen-activated protein kinase genes. ... Mcswiggen J, Beigelman L, Usman N, Haeberli P, Chowrira WPI; 2003-689980/65.

(SIRN-) SIRNA THERAPEUTICS INC.

## Example 3; SEQ ID NO 414; 164pp; English.

The present invention describes a short interfering nucleic acid (siNA) chart downregulates expression of a mitogen-activated protein kinase that downregulates expression of a mitogen-activated protein kinase ((MAPK) genes by RNA interference. Also described: (1) a method for modulating expression of MAPK genes in cells, tissue explants or corganisms by introduction of siNA; (2) kits for in vitro or in vivo delivery of siNA; (3) conjugates and/or complexes of siNA; and (4) vectors that express siNA and cells containing these vectors. MAPK siNAs care cytostatic, anorectic, antidabetic, antihiflammatory.

CC antiathritic, antipsoriatic and gastrointestinal activities. The MAPK siNAs can be used to modulate the expression of MAPK genes, in cells, tissue explants or organisms, e.g. for treating obesity; diabetes types I complish and validation; psoriasis and inflammatory bowel disease. They can also be used for drug screening; diagnosis; target disease. They can also be used for drug screening; diagnosis; target containing agene function and gene mapping (e.g. of single-nucleotide collaboration). The present sequence represents a MAPK sinA which is used in the exemplification of the present invention.

Sequence 19 BP; 4 A; 8 C; 4 G; 0 T; 3 U; 0 Other;

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ö
                          Gaps
                          ;
0
3.0%; Score 12.6; DB 1; Length 19; 68.4%; Pred. No. 4.4e+02; ive 2; Mismatches 4; Indels
                             Conservative
               Local Similarity
                             13;
  Query Match
                Best Loca
Matches
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280 GCGCCACCAAGCTGGTGAA 298 geugecechaleceugha 19

H ADE29888 standard; RNA; 19 (first entry) 29-JAN-2004 ADE29888; ADE29888 RESULT 

Mitogen activated protein kinase siNA oligonucleotide SEQ ID NO:510.

short interfering nucleic acid; siNA; downregulation; inhibition; mitogen-activated protein kinase; MAP kinase; MAPK; RNA interference; cytostatic; anorectic; antidiabetic; antiinflammatory; antiasthmatic; immunosuppressive; antibacterial; antirheumatic; antiarthritic; antipsoriatic; gastrointestinal; obesity; diabetes; tumour; inflammatory disease; asthma; septic shock; rheumatorid arthritis; peoriasis; inflammatory bowel disease; drug screening; genetic engineering; pharmacogenomic; gene mapping; ss.

Synthetic.

WO2003072590-A1.

28-JAN-2003; 2003WO-US002510 04-SEP-2003.

11-MAR-2002; 2002US-0363124P. 06-UUN-2002; 2002US-0386782P. 29-AUG-2002; 2002US-0406784P. 05-SEP-2002; 2002US-0409293P.

(SIRN-) SIRNA THERAPEUTICS INC

15-JAN-2003; 2003US-0440129P

Mcswiggen J, Beigelman L, Usman N, Haeberli P, Chowrira B;

WPI; 2003-689980/65.

New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of mitogen-activated diagnosis of cancer, or protein kinase genes.

Example 3; SEQ ID NO 510; 164pp; English.

The present invention describes a short interfering nucleic acid (siNA) that downregulates expression of a mitogen-activated protein kinase (MAPK) genes by RNA interference. Also described: (1) a method for modulating expression of MAPK genes in cells, tissue explants or organisms by introduction of siNA; (2) kits for in vitro or in vivo organisms by introduction of siNA; (2) kits for in vitro or in vivo corrections that express siNA and cells containing these vectors. MAPK siNAs have cytostatic, anorectic, antidabetic, antifiniammatory, antiasthmatic, immunosuppressive, antibacterial antirheumatic, antipacriatic and gastrointessinal activities. The MAPK siNAs can be used to modulate the expression of MAPK genes, in cells, tissue explants or organisms, e.g. for treating obesity; diabetes types I and II; a wide range of tumours, and inflammatory diseases (asthma, and II; a wide range of tumours, and inflammatory diseases (asthma, disease). They can also be used for drug screening; diagnosis; target cidentification and validation; genetic engineering; pharmacogenomics;

ö studying gene function and gene mapping (e.g. of single-nucleotide polymorphisms). The present sequence represents a MAPK siNA which is used in the exemplification of the present invention. ; 0 3.0%; Score 12.6; DB 1; Length 19; 68.4%; Pred. No. 4.4e+02; ative 2; Mismatches 4; Indels Sequence 19 BP; 3 A; 6 C; 7 G; 0 T; 3 U; 0 Other; 13; Conservative Local Similarity Query Match Best Loc Matches នឧឧឧ

136 CCCGCCTGGCGGTGGAGGC 154

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1 ccueccuesasecuesasec 19

ADE29897 standard; RNA; 19 BP.

ADE29897;

(first entry) 29-JAN-2004 Mitogen activated protein kinase siNA oligonucleotide SEQ ID NO:519

short interfering nucleic acid; siNA; downregulation; inhibition; mitogen-activated protein kinase; MAP kinase; MAPK; RNA interference; cytostatic; anorectic; antidiabetic; antilnitammatory; antiasthmatic; immunosuppessive; antibacterial; antirheumatic; antiarthritic; antipsoriatic; gastrointestinal; obesity; diabetes; tumour; inflammatory disease; asthma; septic shock; rheumatoid arthritis; psoriasis; inflammatory bowel disease; drug screening; genetic engineering; pharmacogenomic; gene mapping; ss

Synthetic.

WO2003072590-A1.

04-SEP-2003

28-JAN-2003; 2003WO-US002510.

11-WAR-2002; 2002US-0363124P. 06-JUN-2002; 2002US-0386782P. 29-AUG-2002; 2002US-0406784P. 05-SEP-2002; 2002US-04092378P. 09-SEP-2003; 2002US-0409293P. 15-JAN-2003; 2003US-0440129P. 

(SIRN-) SIRNA THERAPEUTICS INC.

Mcswiggen J, Beigelman L, Usman N, Haeberli P, Chowrira B;

New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of mitogen-activated protein kinase genes. WPI; 2003-689980/65.

Example 3; SEQ ID NO 519; 164pp; English.

The present invention describes a short interfering nucleic acid (siNA) that downregulates expression of a mitogen-activated protein kinase (MAPK) genes by RNA interference. Also described: (1) a method for modulating expression of NAPK genes in cells, tissue explants or organisms by introduction of SiNA; (2) kits for in vitro or in vivo dellovary of siNA; (3) conjugates and/or complexes of SiNA; and (4) vectors that express siNA and cells containing these vectors. MAPK siNAs have cytostatic, anorectic, antidiabetic, antidiammatory, antiathematic, antidematic, and gastrointestinal activities. The MAPK siNAs antiarthritic, antipeoriatic and gastrointestinal activities. The MAPK siNAs can be used to modulate the expression of MAPK genes, in cells,

tissue explants or organisms, e.g. for treating obesity; diabetes types I and II; a wide range of tumours, and inflammatory diseases (asthma, esptic shock, rheumatorid arthrities, psorifasis and inflammatory bowel disease). They can also be used for drug screening; diagnosis; target identification and validation; genetic enginearing; pharmacogenomics; studying gene function and gene mapping (e.g. of single-nucleotide polymorphisms). The present sequence represents a MAPK sinA which is used in the exemplification of the present invention. 8888888888888

Sequence 19 BP; 3 A; 4 C; 8 G; 0 T; 4 U; 0 Other;

3.0%; Score 12.6; DB 1; Length 19; 78.9%; Pred. No. 4.4e+02; ive 0; Mismatches 4; Indels 280 GCGCCACCAAGCTGGTGAA 298 Local Similarity 78.9 Query Match Matches ઠ

0; Gaps

19 GCTGCCCAACCTGCTGAA 1 원

ADE29783 standard; RNA; 19 29-JAN-2004 (first entry) ADE29783; RESULT 551 ADE29783/C 

short interfering nucleic acid; siNh; downregulation; inhibition; mitogen-activated protein kinase; MAP kinase; MAPK; RNA interference; cytostatic; anridiabetic; antiinflammatory; antiasthmatic; immunosuppressive; antibacterial; antitheumatic; antiarthritic; antipsoriatic; gastrointestinal; obesity; diabetes; tumour; inflammatory disease; asthma; septic shock; rheumatoid arthritis; pesoriasis; inflammatory bowel disease; drug screening; genetic engineering; pharmacogenomic; gene mapping; ss. Mitogen activated protein kinase siNA oligonucleotide SEQ ID NO:405.

Synthetic

WO2003072590-A1.

04-SEP-2003.

28-JAN-2003; 2003WO-US002510.

20-FEB-2002; 2002US-0358580P. 11-MAK-2002; 2002US-0363124P. 06-JUN-2002; 2002US-0386782P. 29-AUG-2002; 2002US-0406784P. 05-SEP-2002; 2002US-0408378P. 09-SEP-2002; 2002US-0409293P. 15-JAN-2003; 2003US-0440129P.

(SIRN-) SIRNA THERAPEUTICS INC.

Mcswiggen J, Beigelman L, Usman N, Haeberli P, Chowrira B;

WPI; 2003-689980/65.

New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of mitogen-activated protein kinase genes.

Example 3; SEQ ID NO 405; 164pp; English.

The present invention describes a short interfering nucleic acid (siNA) that downregulates expression of a mitogen-activated protein Kinase (MAPK) genes by RNA interference. Also described: (1) a method for modulating expression of MAPK genes in cells, tissue explants or organisms by introduction of siNA; (2) kits for in vitro or in vivo delivery of siNA; (3) conjugates and/or complexes of siNA; and (4)

vectors that express sinA and cells containing these vectors. MAPK sinAs have cytostatic, ancrectic, antidiabetic, antinflammatory, anticathmatic, immunosuppressive, antibacterial, anticheumatic, antipacriatic and gastrointestinal activities. The MAPK sinAs can be used to modulate the expression of MAPK genes, in cells, tissue explants or organisms, e.g. for treating obseity; diabetes types I and II; a wide range of tumours, and inflammatory diseases (asthma, septic shock, rheumatoid arthritis, psoriasis and inflammatory bowel disease). They can also be used for drug screening; diagnosis; target dientification and validation; genetic engineering; pharmacogenomics; studying gene function and gene mapping (e.g. of single-nucleotide polymorphisms). The present sequence represents a MAPK sinA which is used in the exemplification of the present invention.

8X8888888888888X8

Sequence 19 BP; 3 A; 7 C; 6 G; 0 T; 3 U; 0 Other;

4, Indels 0, Gaps 3.0%; Score 12.6; DB 1; Length·19; 78.9%; Pred. No. 4.4e+02; Live 0; Mismatches 4; Indels : Conservative Best Local Similarity Matches 15; Conserva Query Match

136 ccccccrccccccqcrccpcc 154 19 CCTGCCTGAAGCTGGAGGC 1 ò 셤

AAZ94278 standard; DNA; 20 BP. AAZ94278

03-JUL-2000 (first entry)

AAZ94278;

PHBLIX; human; testis-specific; transcription factor; prostate cancer; bladder cancer; ovary cancer; testicular cancer; gene therapy; diagnosis; vaccine; PCR primer; ss. Human PHELIX nested primer NP2.

98US-0098610P. 98US-0106524P. 99WO-US020137. WO200012709-A2. 31-AUG-1999; Homo sapiens. 31-AUG-1998; 31-OCT-1998; 09-MAR-2000. 

UROGENESYS INC. AFAR D E. HUBERT R S. RAITANO A B. (HUBE/) (UROG-)

Raitano AB Afar DE, Hubert RS,

WPI; 2000-237872/20.

Testis specific Helix Loop Helix proteins expressed in cancers and useful for the prevention, diagnosis and treatment of prostate, bladder and ovarian tumors.

Example 1; Page 31; 62pp; English.

The present sequence is that of nested primer NP2, which was used in the amplification of gene fragments obtained from a suppression subtractive hybridization reaction using IAPC xenograft cDNA and designed to identify novel prostate and prostate cancer-specific genes. A 437 bp clone was obtained. Full-length cDNA (see AA294275) was subsequently cloned from a transcription factor that is normally expressed only in testis cDNA intervential incommentation factor that is normally expressed only in testis tissue, but is up-regulated in prostate and other types of cancer. The invention

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This sequence represents a PCR primer used in the isolation of cDNA fragments of the PTAN (testis specific protein expressed in prostate cancer) gene. PTAN is expressed in 3 isoforms PTAN-1, 2, and 3. The PTAN gene is located on chromospene 102. PTAN is overexpressed in prostate cancer, and has a testis specific expression pattern in adult tissues. PTAN shows no homology to any known gene. PTAN can be used in methods for the diagnosis of cancer, especially prostate or breast cancer, where the normal tissue samples are prostate itssue, or breast tissue, bone tissue, lymphatic tissue, serum, blood, or urine. A vector containing the PTAN incleotide sequence, a vaccine composition targeting PTAN, PTAN, ribozymes specific for PTAN mRNA and antisense sequences, cancer, especially breast and prostate cancers. Cancer development can be inhibited by a vaccine composition targeting PTAN
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0
provides diagnostic and therapeutic methods useful in the management of various cancers which express PHELIX, including prostate cancer, bladder cancer, ovarian cancer and testicular cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                      PTAN; testis specific; prostate cancer; overexpress; chromosome 1q22; diagnose; cancer; breast; vaccine; PCR primer; ss.
                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PTAN proteins, and sequences encoding them, used for diagnosing and treating cancers, especially breast and prostate cancers.
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                                                                                                Query Match
3.0%; Score 12.6; DB 1; Length 20;
Best Local Similarity 78.9%; Pred. No. 4.9e+02;
Matches 15; Conservative 0; Mismatches 4; Indels
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                                                                     Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mitchell SC;
                                                                                                                                                                                                                                                                                                                                                                                                    PCR primer (NP2) used in PTAN gene isolation.
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                                                                                                                                                                           319 GCGTGCTGGCGGCGACGA 337
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98US-0102910P.
98US-0113229P.
99US-0129518P.
                                                                                                                                                                                                                                                                                               AAA37951 standard; DNA; 20 BP
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HUBERT R S.
RAITANO A B.
MITCHELL S C
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21-DEC-1998;
14-APR-1999;
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BPC-1 polypeptides and polynucleotides can be used for the detection of BPC-1 polypeptides and polynucleotides in biological samples, this is particularly useful for detecting cancers expressing BPC-1, e.g. prostate cancer or bladder cancer. Autibodies directed against BPC-1 or antisense concer or bladder cancer. Autibodies directed against BPC-1 or antisense concerned and be used in vaccines for treating or inhibiting the polymedicetides can also be used for detection, prognosis, drug screening or polymedicetides can also be used for detection, prognosis, drug screening conprises a cTDB domain which is expressed in prostate and bladder comprises a cTDB domain which is expressed in prostate and bladder carcinoma cells and which shows sequence similarity with CTB domains from their known proteins. In normal human tissues BPC-1 is only expressed in certain tissues of the brain, however, it is expressed at high levels in prostate cancer cells and bladder cancer cells. A number of synthetic colligonucleotides were used to generate BPC-1 cDNA from total cell RNA of tumour cells lines. These primars were a cDNA synthesis primer cancer cells in these primars were a cDNA synthesis primer (AAZS3041), two adaptor sequences (AAZS3047, AAZS3048). This sequence is the nested primers (NP)1 used in the amplification method.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New isolated BPC-1 polypeptides, useful for developing products for the diagnosis, staging, prognosis and treatment of cancers, particularly prostate or bladder cancer.
                                                                                                                                                                                                                                                                                                              BPC-1; oncogene; oncogenic; cancer; prostate; bladder; antibody; antisense; vaccine; detection; prognosis; drug screening; primer; ss.
                 Gape
                                                                                                                                                                                                                                                                              Primer used for generating human brain specific protein BPC-1 cDNA.
               ö
Pred. No. 4.9e+02;
0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hubert RS, Leong K, Raitano AB, Saffran DC;
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                                                   319 GCGTGCTGGCGGCGACGA 337
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Best Local Similarity 78.9%;
Matches 15; Conservative
                                                                                                                                                                             AAZ93048 standard; DNA; 20
                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      UROGENESYS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (RAIT/) RAITANO A B.
(SAFF/) SAFFRAN D C.
(JAKO/) JAKOBOVITS A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-206006/18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AFAR D E.
HUBERT R S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LEONG K.
                                                                                                                                                                                                                                                                                                                                                                                                              WO200009691-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Jakobovits A;
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                                                                                                                                                                                                                                                 24-JUL-2000
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                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                               AAZ93048;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (UROG-)
(AFAR/)
(HUBE/)
(LEON/)
                                                                                                                                             RESULT 554
                                                                                                                                                         AAZ93048
ID AAZ9
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3.0%; Score 12.6; DB 1; Length 20;

Query Match

3.0%; Score 12.6; DB 1; Length 20;

Query Match

AAZ94898;

RESULT 555

ò g AAZ94898

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PCR primers AAA14805-07 were used to amplify testis-specific protein Yencoded DNA. The specification describes a new method of diagnosis of specific cancer. The method comprises determining the level of testis-specific protein The method comprises determining the level of testise. TSPY mRNA or protein, and comparing these presence of elevated TSPY mRNA or protein is indicative of prostate cancer. Detection of TSPY mRNA expression or protein levels useful in the diagnosis of prostate cancer. Antisense polymucleotides complementary to the coding sequence of human TSPY are useful for treating prostate cancer by inhibiting TSPY transcription (when contacted with the TSPY mRNA). Ribozymes are also useful for treating prostate cancer by cleaving the TSPY mRNA and therefore inhibiting its translation. The vaccine is useful for the inhibiting the development of prostate cancer in a patient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Diagnosing prostate cancer by determining the level of testis-specific protein Y-encoded (TSPY) mRNA or protein and comparing these TSPY mRNA or protein levels to those of a normal tissue sample.
                                                                                                                                                                                             Prostate cancer, testis-specific protein Y-encoded mRNA, TSPY mRNA, vaccine, PCR primer, 88.
                                                                                                                                                     PCR primer for testis-specific protein Y-encoded DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 1; Page 20; 32pp; English.
                       AAA14807 standard; DNA; 20 BP.
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                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hubert RS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2000-303803/26.
                                                                                                                                                                                                                                                                                                        WO200020638-A2
                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                               02-OCT-1999;
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                                                                 AAA14807;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel testes-specific gene 22P4F11 which is expressed in human prostate cancer and is useful as a diagnostic marker and/or therapeutic target for
                                                                                                                                                                                                                                                                                                                                                                                            22P4F11; human; testis; prostate cancer; diagnosis; gene therapy; marker; vaccine; PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                       Gaps
                                                                                                                                                                                                                                                                                                                                                     PCR primer NP2 used in testis-specific 22P4F11 gene amplification.
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                       4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
Best Local Similarity 78.9%; Pred. No. 4.9e+02; Matches 15; Conservative 0; Mismatches 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hubert RS, Mitchell SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               319 GCGTGCTGGCGGCGACGA 337
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                                                                                                                                                                                                                    AAZ94898 standard; DNA; 20 BP
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99US-0146584P.
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                                                                                                                                                                                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-303452/26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200018925-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-SEP-1999;
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28-JUL-1999;
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3.0%; Score 12.6; DB 1; Length 20;
Best Local Similarity 78.9%; Pred. No. 4.9e+02;
Matches 15; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          36PlA6; transcription factor; murine EHF homologue; ETS cytostatic; cancer; vaccine; tumorigenesis; primer; 9s.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nested primer 2 cloning SSH-generated 36P1A6 gene.
Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                              319 GCGTGCTGGCGGCGACGA 337
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RESULT 556

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99WO-US022576. 98US-0102744P.

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Novel putative transcription factor gene 36P1A6 for treatment, diagnosis and prevention of prostate, bladder, cervical, ovarian, pancreatic, and
                                                                                                            Wer DE, Hubert RS, Mitchell SC;
                                                                                                                                                               Example 1; Page 30; 53pp; English
                                                                            UROG-) UROGENESYS INC.
                                                                                 AFAR D E.
HUBERT R S.
MITCHELL S C.
                                                                                                                        WPI; 2000-303772/26
                                                                                                                                                   colonic cancer.
                  WO200020584-A2.
      Homo sapiens.
                                                         02-0CT-1998;
29-JUL-1999;
                                            02-OCT-1999;
                                13-APR-2000
                                                                                 (AFAR/)
(HUBE/)
(MITC/)
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The human 36PlA6 gene encodes a putative transcription factor based on homology to the murine EHF gene which encodes a transcription factor which is a member of the ETS family. 36PlA6 is expressed in androgen-dependent LAPC prostate cancer xenografts and in normal prostate at approximately equal levels. The highest expression is the the prostate and colon. 36PlA6 may be involved in activating tumor-promoting genes or repressing genes that block tumorigenesis. The 36PlA6 cancer, e.g. prostate, bladder, cervical, ovarian, pancreatic and colonic cancer (all claimed). Anti-36PlA6 antibodies may be used for purifying 36PlA6 and for isolating 36PlA6 antibodies may be used for purifying 36PlA6 and for isolating 36PlA6 antibodies may be used for oligonucleotides and ribozymes can be used to inhibit the transcription and translation of the 36PlA6 gene (Claimed). The 36PlA6 polymucleotides and immunogenic fragments may also be used in cancer and polypeptides and immunogenic fragments may also be used in cancer Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other; (claimed) vaccines

Query Match
3.0%; Score 12.6; DB 1; Length 20;
Best Local Similarity 78.9%; Pred. No. 4.9e+02;
Matches 15; Conservative 0; Mismatches 4; Indels 319 GCGTGCTGGCGGCGGACGA 337

2 degreerededecedada 20

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AAC64567 standard; DNA; 20 BP RESULT 558 AAC64567 

AAC64567;

Human prostate specific 30P3C8 nested primer 2 SEQ ID NO:25.

(first entry)

14-FEB-2001

Human, prostate specific gene, 30P3C8; prostate cancer, diagnosis; cytostatic; gene therapy, vaccine; tumour; primer; 88.

Homo sapiens

WO200061610-A2

19-OCT-2000.

12-APR-2000; 2000WO-US010218

12-APR-1999; 99US-0128860P.

Saffran DC Raitano AB, Leong K, Hubert RS, Afar DE,

WPI; 2000-619224/59 

The present invention describes human prostate specific protein 30P3CB, which is over-expressed in prostate cancer cells. 30P3CB has cytostatic cativity and can be used in vaccines and gene therapy. Methods for detecting the levels of 30P3CB protein or mRNA in prostate tissue, bone tissue, lymphatic tissue, serum, blood or semen are used for diagnoshing the presence of cancer in an individual or disregulated cell growth e.g. hyperplasia. The cancers which are detected or diagnosed are of the bladder, pancreas, colon, brain, bone, lung, kidney or prostate by using test samples of serum, blood or urine or tissues of the bladder, pancreas, colon, brain, bone, lung, kidney and prostate applicable of sequences can be used for treating cancers expressing colymucleotide sequences can be used for treating cancers expressing colymucleotide sequences can be used for treating cancers expressing cused in vaccines to inhibit the development of cancer. Anti-30P3CB are monoclonal antibodies bind to 30P3CB and disrupt interactions between condense and other proteins e.g. receptors for which 30P3CB is a ligand. 30P3CB may be a growth factor or other molecule involved in tumour growth cancer promoting activities of 30P3CB. The assays are invasion or other cancer promoting activities of 30P3CB. The assays are used for detecting, staging and monitoring prostate cancer. The 30P3CB cused for detecting provide a more specific assay than the serum prostate concer and provide a more specific assay than the serum prostate concer and provide a more specific assay than the serum prostent energy present appreaent sequence represents a 30P3CB conserted primer, which is used in the exemplification of the present

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

Gaps .. 0 Query Match
3.0%; Score 12.6; DB 1; Length 20;
Best Local Similarity 78.9%; Pred. No. 4.9e+02;
Matches 15; Conservative 0; Mismatches 4; Indels

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Gaps

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AAC64486;

13-FEB-2001 (first entry)

Prostate tumour associated gene 24P4C12 nested primer 2 SEQ ID NO:41. 

Human; prostate tumour associated gene; 24P4C12; prostate cancer; transmembrane protein; diagnosis; anticancer; cytostatic; vaccine; gene therapy; PCR primer; ss.

(UROG-) UROGENESYS INC

30P3C8 polypeptide and polynucleotide used for diagnosing, treating and monitoring development of prostate cancer.

Example 1; Page 57; 99pp; English.

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RESULT 559 AAC64486

AAC64486 standard; DNA; 20 BP.

Homo sapiens.

WO200061746-A1.

19-OCT-2000.

12-APR-2000; 2000WO-US010039

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Novel 24P4C12 polypeptides and polynucleotides, used in the diagnosis and treatment of cancer, especially prostate cancer.
                                                                                                                                                                                                                                                                                       The present invention describes a prostate tumour associated gene, designated 24P4C12, and its encoded protein. 24P4C12 has anticancer and cytostatic activity, and can be used in vaccine production and in gene therapy. A pharmaceutical composition or vaccine comprising 24P4C12 can be used to treat a patient with cancer, especially prostate cancer, the vaccine can also be used to inhibit the development or progression of cancer. The polypeptides and polynucleotides can be used to disgnose cancers, especially prostate cancer. A transgenic animal comprising 24P4C12 can be used for the development and screening of therapeutic reagents. The polypeptide is a transmembrane protein which is expressed specifically in prostate cancer, allowing the development of more specific anticancer therapies and diagnostic assays
                                                                                        Leong K, Raitano AB, Saffran DC;
                                                                                                                                                                                                                                                   Example 1; Page 65; 137pp; English.
  12-APR-1999; 99US-0128858P.
                                         (UROG-) UROGENESYS INC.
                                                                                        Hubert RS,
                                                                                                                                    WPI; 2000-672681/65.
                                                                                        Afar DE,
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0; Gaps 3.0%; Score 12.6; DB 1; Length 20; 78.9%; Pred. No. 4.9e+02; tive 0; Mismatches 4; Indels Local Similarity 78.9 Query Match

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

AAF85709 standard; DNA; 20 BP 10-DEC-2001 (first entry) AAF85709; RESULT 560 

Human, cancer related protein 20P2H8; vaccine; chromosome 15q32-23; prostate cancer; bladder cancer; colon cancer; pancreatic cancer; PCR primer; ss. Human cancer related protein 20P2H8 cDNA PCR primer #3.

Homo sapiens.

WO200131012-A1.

03-MAY-2001

26-OCT-2000; 2000WO-US029477. 99US-0162364P. 28-OCT-1999;

(UROG-) UROGENESYS INC.

Hubert RS, Mitchell SC, Jakobovits A; Afar DEH, Raitano AB, WPI; 2001-308645/32. 20P2H8 polynucleotides and polypeptides useful for diagnosing and treating cancer, and for screening for screening for modulating treating c

Example 1; Page 64; 111pp; English.

ö The present invention provides the protein and coding sequences of human cancer related protein 20P2H8. The gene, which is found at chromosome 15q32-23, is upregulated in cancers such as that of the prostate, bladder, colon and pancreas. The sequences can be used to diagnose and treat these cancers, and to vaccinate against them. The present sequence is a PCR primer for the coding sequence of the invention Gaps . 0 3.0%; Score 12.6; DB 1; Length 20; 78.9%; Pred. No. 4.9e+02; tive 0; Mismatches 4; Indels Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other; 319 GCGTGCTGGCGGCGGACGA 337 Query Match
Best Local Similarity 78.9
Matches 15; Conservative

8888888

2 GCGTCGCGCCCGAGGA 20 ਨੇ 셤

AAD06232 standard; DNA; 20 BP.

RESULT 561

AAD06232; 

(first entry) 31-JUL-2001

Human SGP28 gene fragment amplifying NP2 primer.

Human, specific granule protein 28; SGP28; therapy; PCR primer; prostate; colon; cancer; prognosis; vaccine; anticancer; SSH; suppression subtractive hybridisation; ss.

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03-MAY-2001.

WO200131343-A2.

27-OCT-2000; 2000WO-US029607.

99US-0162610P. 28-OCT-1999;

(UROG-) UROGENESYS INC.

Raitano AB, Afar DEH, Mitchell SC, Faris M; Jakobovits A; Hubert RS,

WPI; 2001-308685/32.

Detecting cancers, particularly of prostate and colon, from overexpression of SGP28 protein, also methods for treating these cancers e.g. by vaccination with the protein.

Example 1; Page 59; 102pp; English.

The present invention relates to methods and compositions for the diagnosis and therapy of prostate cancer which utilise human SGP28 (specific granule protein 28) gene and proteins. The method involves ceptering cancers, particularly of prostate and colon, from cancers, particularly of prostate and colon, from coveragesion of SGP28 protein. The expression of SGP28, which is an coveracelular protein is restricted to the prostate and ovary, and is markedly up-regulated in prostate tumours. SGP28 sequence is used for diagnosis (including in vivo inaging), staging, monitoring and prognosis of prostatic and colon cancer, and for assisting selection of therapy. Composition or vaccine that contains a vector expressing an antibody specific for SGP28 protein, nucleic acid encoding SGP28 protein or the specific antibody opionally conjugated to toxin or therapeutic agent. SGP28 gene product is also used as source of therapeutic antisense or ribosyme agents, as primers/probes for diagnosis or prognosis, to identify compounds that inhibit calcium entry into prostatic cells, for recombinant production of SGP28 peptides and for isolating related sequences. SGP28 protein and its

Wed Apr 21 12:58:21 2004

fragments are used to raise specific antibodies (Ab) and to identify specific binding agents (potentially useful as therapeutic and diagnostic agents) and also potential anticancer agents. The present sequence is a nested primer 2 (NP2) used to amplify gene fragments resulting from SSH (suppression subtractive hybridisation) reaction. This sequence is used in the SSH isolation of cDNA fragment of human SGP28 gene 8888888888

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

0; Gaps 3.0%; Score 12.6; DB 1; Length 20; ilarity 78.9%; Pred. No. 4.9e+02; Conservative 0; Mismatches 4; Indels Local Similarity nes 15; Conserva Query Match Best Loca Matches

RESULT 562 AAD04811

AAD04811 standard; DNA; 20 BP. AAD04811; 

(first entry) 17-JUL-2001

Human 36P6D5 gene fragment amplifying primer NP2.

Human, 16P6D5 protein; secreted tumour antigen; therapy; cancer; kidney; bladder; ovary; breast; pancreas; colon; lung; vaccine; cytostatic; SSH; suppression subtractive hybridisation; PCR primer; ss.

Homo sapiens

WO200131015-A2.

30-OCT-2000; 2000WO-US029894.

99US-0162417P. 28-OCT-1999;

(UROG-) UROGENESYS INC

Jakobovits A, Faris M, Afar DEH, Hubert RS; Raitano AB, Mitchell SC;

WPI; 2001-308646/32.

Detecting presence of cancer expressing 36P6D5 protein in individual by comparing protein level in test sample to normal sample, where elevated level of protein in test sample indicates presence of cancer.

Example 1; Page 70; 113pp; English.

The present invention relates to a gene and its encoded secreted tumour antigen, termed 36P6D5. These sequences are used for the diagnosis and treatment of various cancers which express 36P6D5, such as cancers of the kidney, bladder, ovary, breast, pancreas, colon and lungs. In normal individuals 36P6D5 protein, is predominantly expressed in pancreas, with comparising immunogenic protein of 36P6D5 is useful for inhibiting the development of prostate or colon cancer. Pharmaceutical composition comprising 36P6D5 protein is useful for diagnosis and/or programs of prostate cancer and other cancers, for modulating or inhibiting the expression of 36P6D5 genes and/or translation of the 36P6D5 transcripts, and as therapeutic agents. The present sequence is a nested primer (NP)2 used to amplify gene fragments resulting from SSH (suppression subtractive hybridisation) reaction. This sequence is used in the SSH isolation of cDNA fragment of human 36P6D5 gene

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

Gaps ö 3.0%; Score 12.6; DB 1; Length 20; 78.9%; Pred. No. 4.9e+02; ive 0; Mismatches 4; Indel8 Query Match Best Local Similarity 78.9 Matches 15; Conservative

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319 GCGTGCTGGCGGCGACGA 337 2 degredecedecedada 20 g ઠ

AAF76012 standard; DNA; 20 BP

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AAF76012;

22-MAY-2001

PCR primer NP2, SEQ ID NO:18, used in human PC-LECTIN cDNA isolation.

Human; PC-LECTIN; C-type lectin; transmembrane antigen; normal testis; layilin homologue; prostate cancer antigen; overexpression; androgen-dependent prostate cancer; diagnosis; prognosis; PCR primer; ss.

Synthetic.

WO200112811-A1.

22-FEB-2001

11-AUG-2000; 2000WO-US022065.

99US-0148935P. 12-AUG-1999;

(UROG-) UROGENESYS INC

Jakobovits A, Raitano AB; Afar DEH, Hubert RS,

WPI; 2001-211222/21.

New PC-LECTIN polynucleotide encoding a transmembrane antigen over expressed in human prostate cancer, useful for the prognosis, diagnosis and treatment of prostate cancer.

Example 1; Page 59; 116pp; English.

The invention relates to a novel human C-type lectin transmembrane cantigen, PC-LECTIN (AAB73309) and cDNA encoding it (AAF76004). The antigen, PC-LECTIN (AAB73309) and cDNA encoding it (AAF76004). The capression of the human PC-LECTIN gene is normally restricted to the captesion is highly overexpressed in prostate cancer. PC-LECTIN therefore independent prostate tumours, and expression is therefore likely conditional dependent prostate tumours compared with androgen-independent prostate tumours. And expression is therefore likely cropsession is the presence of androgen. Human PC-LECTIN therefore cropsession adiagnostic and therapeutic target for prostate cancer. Cropsession androgen-dependent prostate cancer. Human PC-LECTIN exhibits concilorly to hamster layilin (44.9% identity over a 265 residue overlap), concilorly in a key functional domain proposed for the layilin concilor protein. Human PC-LECTIN applicational domain proposed for the layilin concilor protein. Human PC-LECTIN antisense nucleotide, a PC-LECTIN concilor-treating a patient with a cancer, concilor-targetted ribozyme, or an anti-PC-LECTIN antibody may be used to prepare a composition for treating a patient with a cancer, colon, pancreatic, testicular, cervical or ovarian cancers that express colon, pancreatic, testicular, cervical or ovarian cancers that cxpress colon, pancreatic, testicular, cervical or ovarian cancers for colon, pancreatic, testicular, cervical or ovarian cancers for colon, pancreatic, testicular, each also useful for diagnosing the presence of cancer. PC-LECTIN antibodies and nucleotides are useful in the antibodies may also be used as drug targetting agents. The PC-LECTIN antibodies may also be used as drug targetting agents. The PC-LECTIN concludes and proteins may additionally be used in the isolation of human concern sequence represents a pCR-LECTIN function or expression. 

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Gaps .; 0 3.0%; Score 12.6; DB 1; Length 20; llarity 78.9%; Pred. No. 4.9e+02; Conservative 0; Mismatches 4; Indels Local Similarity hes 15; Conserv Query Match Matches

AAF83890 standard; DNA; 20 BP. RESULT 564 AAF8389(

06-AUG-2001 (first entry)

Nested primer (NP)2 used in human PHOR-1 cDNA isolation.

G-protein-coupled receptor; prostate; cancer; PHOR-1; kidney; uterine; cervical; stomach; rectal; cytostatic; vaccine; cell function regulator; human; prostate homologue of olfactory receptor-1; PCR primer; se.

Homo sapiens.

WO200125434-A1.

12-APR-2001

05-OCT-2000; 2000WO-US027543.

05-OCT-1999;

(UROG-) UROGENESYS INC.

Jakobovits A, Faris M, Hubert RS; Afar DEH, Ja , Saffran DC; Raitano AB, Mitchell SC,

WPI; 2001-367230/38.

Novel gene designated PHOR-1, a G-protein-coupled receptor up-regulated in prostate cancer, useful as diagnostic marker and therapeutic target for cancers of prostate, kidney, uterus.

Example 1; Page 59; 139pp; English.

The invention relates to a novel G-protein-coupled receptor up-regulated in prostate cancer, termed PHOR-1. The encoding cDNA is contained in plantal designated plo1019All deposited with ATC as Accession No.PTA-112. PHOR-1 polypepides and polymuclectides are useful for diagnosing the presence of cancer, especially prostate, kidney, uterine, cervical, cancer where a comment is a comparing the level of the protein or mRNA expression in test and comparing the level of the protein or mRNA compositions comprising PHOR-1 is useful for treating cancer. PHOR-1 proteins are useful for identifying ligands and other of generating antibodies which are useful in diagnostic, prognostic and for generating methodologies and for the treatment of prostate cancer. Cell insign methodologies and for the treatment of prostate cancer. Cell interactions mediated by PHOR-1 (The present sequence represents a primer useful in isolation of the PHOR-1 (prostate homologue of olfactory receptor

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

Gaps ò Query Match 3.0%; Score 12.6; DB 1; Length 20; Best Local Similarity 78.9%; Pred. No. 4.9e+02; Matches 15; Conservative 0; Mismatches 4; Indels

319 GCGTGCTGGCGGCGGACGA 337

2 GCGTGGTCGCGGCCGAGGA 20

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RESULT 565 AAH99163

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AAH99163 standard; DNA; 20 BP.

AAH99163;

(first entry) 04-DEC-2001 Human prostate-related gene 83P5G4 cDNA nested primer #2.

83P5G4; PCR primer; DNA adaptor; prostate; testis; tissue; cancer; ss; tumour; kidney; brain; bone; ovary; breast; pancreas; uterus; colon; lung; cytostatic; gene therapy; antibody therapy; ribozyme; liver; single chain monoclonal antibody; serum; blood; urine; bladder; cervix; rectum; stomach; human; chromosome 1q31-q32.

Homo sapiens.

WO200159115-A2.

16-AUG-2001.

09-FEB-2001; 2001WO-US004426.

09-FEB-2000; 2000US-0181261P. 

(UROG-) UROGENESYS INC.

Challita-Eid PM, Faris M, Levin E;

An isolated 83P5G4-related protein useful as a diagnostic and/or therapeutic agent in multiple cancers such as prostate, bladder and bone

Example 1; Page 55; 112pp; English.

The nucleic acid sequences represent the 83P5G4 gene and the primers and adaptors used to amplify 83P5G4 DNA. 83P5G4 exhibits prostate specific expression in normal adult tissue, but it is also aberrantly expressed in many cancers including tumours of the prostate, testis, bladder, kidney, brain, bone, cervix, uterus, ovary, breast, pancreas, stomach, rectum, liver, colon and lung The 83P5G4 polynucleotide, its related protein and also peptide fragments of the protein are therefore useful for diagnosing and treating cancer. A vector comprising a polynucleotide which encodes a single chain monoclonal antibody, that immunospecifically binds to an single chain monoclonal antibody, that immunospecifically binds to an plynucleotide having the 83P5G4 capable of cleaving; a polynucleotide having the 83P5G4 capable of cleaving; a persention of a composition for treating a patient with a cancer that expresses 83P5G4. The sequences can be used in diagnostic methods to monitor the level of 83P5G4 gene products in serum, blood, urine and to thereby detect the presence of cancerous cells

ö Query Match
3.0%; Score 12.6; DB 1; Length 20;
Best Local Similarity 78.9%; Pred. No. 4.9e+02;
Matches 15; Conservative 0; Mismatches 4; Indels;

Gaps

2 dédrégredédededada 20

AAS42202 standard; DNA; 20 BP.

Jakobovits A; Hubert RS, Afar DEH, Mitchell SC, Jakobovit

WPI; 2001-514669/56.

cancer.

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

319 GCGTGCTGGCGGCGACGA 337

RESULT 566 AAS42202 ID AAS4220 XX

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prostate; colon; bladder; lung; ovarian; pancreatic; PCR primer; ss
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Best Local Similarity 78.9
Matches 15; Conservative
                                                                                                                                                                                                                                                         Hubert RS, F
Jakobovits A;
                                                                                                                                                                                                                      (UROG-) UROGENESYS INC
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                                                                        WO200140276-A2
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                                       Homo sapiens.
                                                                                                           07-JUN-2001.
                                                                                                                                                                                                                                                         Afar DEH,
Faris M,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequences AAS42193-AAS42208 represent the 103P2D6 gene and the primers and adaptors used to amplify 103P2D6 DNA. 103P2D6 is not expressed in normal adult tissue but is aberrantly expressed in some foretal tissues contain and also protects, testis, bladder, bone, cervix, ovary, breast, pancreas, colon and lung. The 103P2D6 DOPE, comprising a polymucleotide protein and also peptide fragments of the protein are therefore useful for diagnosing and treating cancer. A vector comprising a polymucleotide which encodes a single chain monoclonal antibody, that immunospecifically binds to an 103P2D6-related protein, antibody, that immunospecifically binds to an 103P2D6-related protein, and a ribozyme capable of cleaving a polymucleotide having the 103P2D6 coding sequence, are both useful in the preparation of a composition for treating a patient with a cancer that expresses 103P2D6. The sequences can be used in diagnostic methods to monitor the level of 103P2D6 gene products in serum, blood, urine and tissue and to thereby detect the presence of cancerous cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human, cytostatic, antiproliferative, vaccine, gene therapy,
six transmembrane epithelial antigen of the prostate-1; STEAP-1; cancer;
                                                                                                       103P2D6; PCR primer; DNA adaptor; prostate; testis; foetal tissue; 8s; tumour; cancer; bone; ovary; breast; pancreas; colon; lung; cytostatic; gene therapy; antibody therapy; ribozyme; serum; blood; urine; bladder; single chain monoclonal antibody; cervix; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NP2 primer used in isolation of STEAP cDNA fragment generated from SSH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New polynucleotide for treating and diagnosing prostate cancer is the 103P2D6 gene which encodes for 103P2D6-related proteins.
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Raitano AB, Afar DEH, Rastegar GS, Mitchell SC, Hubert RS;
Challita-Eid PM, Faris M, Jakobovits A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.0%; Score 12.6; DB 1; Length 20; 78.9%; Pred. No. 4.9e+02; tive 0; Mismatches 4; Indels
                                                                        fuman prostate-related gene 103P2D6 cDNA nested primer #2.
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                                       (first entry)
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Best Local Similarity 78.5.
Best Local Si Conservative
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                                       17-DEC-2001
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 AAS42202;
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AAD07091
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the present sequence is nested primer (NP2) which is used to isolate the human six transmembrane epithelial antigen of the prostate (STEAP) cDNA fragment generated from suppression subtractive hybridisation (SSH).

STEAP gene is used in gene therapy. Inhibiting the development or progression of a cancer (eg. prostate, colon, bladder, lung, ovarian and pancreatic) expressing STEAP or inhibiting powth or killing cells capressing STEAP, or inhibiting the development or composition to the patient. Treating a patient with a cancer that expresses STEAP, or inhibiting growth or killing cells expressing STEAP, comprises administering to the patient of the patient of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New STEAP (six transmembrane epithelial antigen of the prostate) proteins, expressed in human cancers, useful for detecting and treating
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                                                                                                                                                                                                                                                                                                                                                                                                          Raitano AB, Saffran DC, Mitchell SC;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 70; 187pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    319 GCGTGCTGGCGGCGACGA 337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2 dégrégregédegédegá 20
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06-DEC-2000; 2000WO-US033040.
                                                                                                                                     99US-00455486
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Page 274

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Wed Apr 21 12:58:21 2004
                        26-JAN-2001; 2001WO-US002651.
                                26-JAN-2000; 2000US-0178560P.
                                               Jakobovits A, Afar DEH,
Hubert RS;
                                       (UROG-) UROGENESYS INC
                                                          WPI; 2001-502631/55.
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Challita-Eid PM, Levin E, Mitchell SC;

The nucleic acid sequences represent the 84P2A9 gene and the primers and adaptors used to amplify 84P2A9 DNA. 84P2A9 exhibits prostate and testis specific expression in normal adult tissue, but it is also aberrantly expressed in many cancers including leukaemia and tumours of the prostate, testis, kidney, brain, bone, skin, ovary, breast, pancreas, colon and lung. The 84P2A9 polymuclecide, its related protein and also peptide fragments of the protein are therefore useful for diagnosing and treating cancer. A vector comprising a polymuclectide which encodes a single chain monoclonal antibody, that immunospecifically binds to an 84P2A9-related protein, and a ribozyme capable of cleaving a polymuclectide having the 84P2A9 coding sequence, are both useful in the preparation of a composition for treating a patient with a cancer that expresses 84P2A9. The sequences can be used in diagnostic methods to monitor the level of 84P2A9 gene products in serum, blood, urine and tissue and to thereby detect the presence of cancerous cells New 84P2A9 gene and its encoded protein, useful for diagnosing and treating cancer, e.g. leukemia and cancer of the prostate, testis, kidney, brain or bone, or for eliciting an immune response. Sxample 1; Page 71; 149pp; English.

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

4; Indels 0; Gaps Query Match 3.0%; Score 12.6; DB 1; Length 20; Best Local Similarity 78.9%; Pred. No. 4.9e+02; Matches 15; Conservative 0; Mismatches 4; Indels

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ABL50419 standard; DNA; 20 BP 17-JUN-2002 (first entry) ABL50419; RESULT 569 ABL50419 

Human 158P1F4 gene nested primer (NP)2 SEQ ID NO:736.

Human; 158P1H4; chromosome 8q220q23, 158P1F4; chromosome 8q23; cancer; bladder cancer; immune response; cytotoxic T lymphocyte; CTL; HLA; human leukocyte antigen; helper T lymphocyte; HTL; PCR primer; adapter;

sapiens Synthetic. Homo

WO200216598-A2.

28-FEB-2002.

22-AUG-2001; 2001WO-US026411.

22-AUG-2000, 2000US-0227098P. 10-APR-2001; 2001US-0282739P.

(AGEN-) AGENSYS INC

Levin E; Raitano AB, Afar DEH, Challita-Eid PM, Hubert RS, F Faris M, Ge W, Jakobovits A;

WPI; 2002-269357/31.

Monitoring 158P1H4 gene products in biological sample from patient who has or is suspected of having cancer, useful for treating cancer, comprises identifying presence of aberrant 158P1H4 gene products in biological sample.

Example 45; Page 116; 209pp; English.

The present invention describes a method for monitoring 158P1H4 gene

Comparison a biological sample from a patient who has or is suspected of

Eximp cancer. The method comparises determining the status of 158P1H4

Gene products in a tissue sample from an individual, comparing the status

Co the status of 158P1H4 gene products in a normal sample, and

Co identifying the presence of aberrant 158P1H4 gene products in the sample.

Co identifying the presence of aberrant 158P1H4 gene products in the sample.

Co production. 158P1H4 polynucleotides may be used in monitoring genetic

abnormalities. The 158P1H4 proteins may be used in most genetic

co insepl1H4 gene products in normal versus cancerous tissues and consin-specific antibodies, for identifying agents or cellular factors

Co insepl1H4 general phenotype, in generating and characterising

co lissp1H4 or its particular domain, and for generating cancer

co that bind to 158P1H4 or its particular domain, and for generating cancer

co that bind to 158P1H4 are useful in diagnostic and

concines. Antibodies against 158P1H4 are useful in diagnostic and

concines. Antibodies are particularly useful in bladder cancer diagnostic and

antibodies are particularly useful in bladder cancer diagnostic and

concined in memological reagents for detecting 158P1H4 expressing cells. The

concined to chromosome 8422-q23, and the 158P1H4 gene also described in

the present invention has been located to chromosome 8q23. ABL50400 to

ABL50425 and ABB94468 to ABB95188 represent sequences used in the exemplification of the present invention

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

Gaps 4; Indels 0; Query Match
3.0%; Score 12.6; DB 1; Length 20;
Best Local Similarity 78.9%; Pred. No. 4.9e+02;
Matches 15; Conservative 0; Mismatches 4; Indels

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319 GCGTGCTGGCGGCGGACGA 337 2 dcdrdgrcdcddccdagda 20 ð g

ABL50407 standard; DNA; 20 BP. 17-JUN-2002 (first entry) ABL50407; RESULT 570 ABL50407

Human 158P1H4 gene nested primer (NP)2 SEQ ID NO:724.

Human; 158P1H4; chromosome 8q220q23, 158P1F4; chromosome 8q23; cancer; bladder cancer; immune response; cytotoxic T lymphocyte; CTL; HLA; human leukocyte antigen; helper T lymphocyte; HTL; PCR primer; adapter;

Homo sapiens.

Synthetic.

28-FEB-2002

22-AUG-2001; 2001WO-US026411.

22-AUG-2000; 2000US-0227098P.

rng.res

WO200216598-A2. 

10-APR-2001; 2001US-0282739P

(AGEN-) AGENSYS INC.

Levin E; Raitano AB, Afar DEH, Challita-Eid PM, Hubert RS, 1 Faris M, Ge W, Jakobovits A;

VPI; 2002-269357/31

Monitoring 158PIH4 gene products in biological sample from patient who has or is suspected of having cancer, useful for treating cancer, comprises identifying presence of aberrant 158PIH4 gene products in biological sample.

Example 1; Page 69; 209pp; English

The present invention describes a method for monitoring 158P1H4 gene

products in a biological sample from a patient who has or is suspected of

baving cancer. The method compises determining the status of 158P1H4

gene products in a tissue sample from an individual, comparing the status

contained the presence of aberrant 158P1H4 gene products in the sample.

Contained the presence of aberrant 158P1H4 gene products in the sample.

Contained the presence of aberrant 158P1H4 gene products in the sample.

Contained the presence of aberrant 158P1H4 gene products in the sample.

Contained the presence of a proteins may be used in monitoring genetic

abnormalities. The 158P1H4 proteins may be used in monitoring genetic

contained the malignant phenotype, in generating and characterising

contained the patricular domain, and for generating and contained proteins against 158P1H4 are useful in diagnostic and

contained to the patricular domain, and for generating

contoined a particularly useful in bladder cancer in generating

contoined assays, in treating patients with cancer, in generating

contoined assays, and imaging methodologies. The 158P1H4 gene has been

contibodies are particularly useful in bladder cancer diagnostic and

prognostic assays, and imaging methodologies. The 158P1H4 gene has been

contibodies are particularly useful in bladder cancer diagnostic and

prognostic assays, and imaging methodologies. The 158P1H4 gene has been

content invention has been located to chromosome 8q23. ABL50400 to

ABL50429 and ABB9468 to ABB95188 represent sequences used in the

content of the present invention

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

0; Gaps Ouery Match 3.0%; Score 12.6; DB 1; Length 20; Best Local Similarity 78.9%; Pred. No. 4.9e+02; Matches 15; Conservative 0; Mismatches 4; Indels

319 gegreeregeegeacea 337 śceredrececeska 20

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RESULT 571

ABA98342 standard; DNA; 20 ABA98342; ABA98342 

29-NOV-2002 (first entry)

Nested primer (NP) 2.

55P4H4; cancer; immune response; ds; PCR primer.

Jnidentified.

20-DEC-2001.

13-JUN-2001; 2001WO-US019246

13-JUN-2000; 2000US-0211454P

(UROG-) UROGENESYS INC.

Levin E, Mitchell SC, Raitano AB; Faris M, Hubert RS, Afar DEH, Jakobovits A;

WPI; 2002-098053/13.

Novel isolated 55P4H4-related protein encoded by a gene over-expressed in multiple cancers, useful as a diagnostic and/or therapeutic agent for cancer, preferably prostate cancer.

Example 1; Page 54; 160pp; English

This invention relates to an isolated 55P4H4-related protein encoded by a conservation relates to an isolated 55P4H4 related protein, providing the protein comprises of at least one T cell or B cell epitope. The immune system cell is a B cell which generates antibodies that comprises of at least one T cell or B cell epitope. The immune system cell is a B cell which generates antibodies that comprises of autologous cell that expresses the 55P4H4 controlled which kills an autologous cell that expresses the 55P4H4 controlled which kills an autologous cell that expresses the 55P4H4 controlled which is considered useful for monitoring the presence of facilitate the cytotoxic activity of a cytotoxic T lymphocyte. A method is mentioned which is considered useful for monitoring the presence of cancer in an individual, where the presence of elevated 55P4H4 mRNA or protein expression in the test sample relative to the normal tissue concurs in a prostate, kidney, testis, lung cervix, bone, bladder, brain or ovary tissue. The protein is useful in diagnostic assays that examine contituons associated with disregulated cell growth such as cancer and is also useful in foremaic analysis of tissues of unknown origin, to treat a pathological condition characterized by the overexpression of 55P4H4, for assessing the status of 55P4H4 gene products in normal versus cancer use cissues and to assess the presence of perturbations in specific regions of the 55P4H4 gene. This sequence represents nested primer (NP) 2 used curing the method highlighted in the examples

Sequence 20 BP; 3 A; S C; 10 G; 2 T; 0 U; 0 Other;

Query Match
Best Local Similarity 78.9%; Pred. No. 4.9e+02;
Matches 15; Conservative 0; Mismatches 4; Indels

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RESULT 572 ABA03609

ABA03609 standard; DNA; 20 BP.

ABA03609;

08-FEB-2002 (first entry)

Nested primer 2 used for human 34P3D7 cDNA isolation.

Human, 34P3D7; cytostatic; vaccine; gene therapy; cancer; human leukocyte antigen; HLA; major histocompatibility complex; MHC; HLA A1; HLA A11; HLA A02; HLA A24; HLA A3; HLA B35; HLA B7; primer; 88. 

Homo sapiens

WO200159110-A2.

16-AUG-2001.

08-FEB-2001; 2001WO-US004094

08-FEB-2000; 2000US-0181020P.

(UROG-) UROGENESYS INC.

The invention relates to a polynucleotide, designated 3473D7, encoding 3473D7.related protein, comprising a sequence of 2198 nucleotides fully defined in the specification. The presence of elevated 3473D7 mRNA or protein expression indicates the presence of cancer occurring in protein expression indicates the presence of cancer occurring in breast, pancreatic, ridmey, brain, bone, cervical, uterine, ovarian, breast, pancreatic, stomach, colon, rectal leukocytes, liver, and lung protein, an antisense polynucleotide complementary to 3473D7 related polynucleotide, or a ribozyme capable of cleaving the 34P3D7 polynucleotide, are aribozyme capable of cleaving the 34P3D7 polynucleotide is useful for inhibiting the development of a cancer expressing 34P3D7 in a patient. The present sequence was used in an example demonstrating suppression subbractive hybridisation (SSH)-generated isolation of a cDNA fragment of the 34P3D7 gene New gene designated 34P3D7, encoding a tissue-specific protein highly expressed in prostate cancer, for use as diagnostic and/or therapeutic target for cancers, and for eliciting an immune response. Faris M, Afar DEH, Challita-Bid PM, Hubert RS, Levin E; Mitchell SC, Jakobovits A; 3.0%; Score 12.6; DB 1; Length 20; 78.9%; Pred. No. 4.9e+02; Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other; 0; Mismatches Example 1; Page 53; 112pp; English. 319 GCGTGCTGGCGGCGGACGA 337 2 dcéregrededecedades 20 Best Local Similarity 78.9 Matches 15; Conservative WPI; 2002-025689/03 

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Human 125P5C8 gene PCR primer #3. AALS0002 standard; DNA; 20 BP 10-DEC-2002 (first entry) AAL50002; RESULT 573 

Human, 125P5C8, cancer, cytostatic, breast cancer, prostate cancer, bladder cancer, kidney cancer, colon cancer, ovarian cancer, PCR, primer,

Homo sapiens.

WO200272785-A2.

19-SEP-2002.

14-MAR-2001; 2001US-00809638

13-MAR-2002; 2002WO-US007855.

(AGEN-) AGENSYS INC.

Ge ₩; Afar DEH, Raitano AB, Faris M, Challita-Eid PM, Hubert RS, P Morrison RK, Morrison K, Jakobovits A;

WPI; 2002-713510/77.

New composition comprising a substance that modulates the status of 125P5C8 gene or a molecule that is modulated by 125P5C8, useful for treating or preventing cancer that expresses or over expresses 125P5C8.

Example 1; Page 68; 274pp; English

The present invention relates to compositions comprising a substance that modulates the status of 125P5C8 or a molecule that is modulated by 125P5C8. The status of a cell that expresses 125P5C8 is modulated by composition is useful for treating cancer, particularly prostate, bladder, kidney, colon, ovary or breast cancer. The 125P5C8 protein amount of a nucleotide sequence encoding the protein is useful for immunising a mammal against cancer. The present sequence is a PCR primer shown in the exemplification of the invention . Match 3.0%; Score 12.6; DB 1; Length 20; Local Similarity 78.9%; Pred. No. 4.9e+02; es 15; Conservative 0; Mismatches 4; Indels Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other; 319 GCGTGCTGGCGGCGACGA 337 2 decirectedecedada 20 AAS95820 standard; DNA; 20 BP. Query Match Best Loca Matches 8888888888888 d

103F9B8; PCR primer; DNA adaptor; prostate; bladder; kidney; colon; lung; breast; rectum; stomach; tumour; cancer; cytostatic; gene therapy; ss; antibody therapy; ribozyme; single chain monoclonal antibody; serum; blood; urine; tissue; human; chromosome 9q13-q21. Human cancer-related gene 103P3E8 cDNA nested primer #2.

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0; Gaps

4; Indels

26-FEB-2002 (first entry)

WO200179557-A2.

25-OCT-2001

12-APR-2001; 2001WO-US012181.

12-APR-2000; 2000US-0196647P.

(UROG-) UROGENESYS INC

Challita-Eid PM, Raitano AB, Mitchell SC, Afar DEH; Jakobovits A; Paris M,

NPI; 2002-061976/08.

Monitoring 103P3E8 gene products in sample from patient (suspected of) having cancer, useful for diagnosing, managing or treating cancers, e.g. prostate cancer, comprises determining presence of aberrant 103P3E8 gene 

Example 1, Page 55, 128pp; English.

Sequences AAS95810-AAS95820 represent the 1039388 gene and the primers and adaptors used to amplify 1039388 DNA. 1039388 exhibits tissue specific expression in normal adult tissue, but it is also aberrantly expressed in many cancers including tumours of the prostate, bladder, kidney, colon, lung, breast, rectum and stomach. The 103938 polynucleotide, its related protein and also peptide fragments of the protein are therefore useful for diagnosing and treating cancer. A vector comprising a polynucleotide which encodes a single chain monoclonal antibody, that immunospecifically binds to an 1039388-related protein, and a ribozyme capable of cleaving a polynucleotide having the 1039388 coding sequence, are both useful in the preparation of a composition for treating a patient with a cancer that expresses 1039388. The sequences can be used in diagnostic methods to monitor the level of 1039388 gene products in serum, blood, urine and tissue and to thereby detect the

presence of cancerous cells ន្តដូន

Seguence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

Gaps ö Length 20; 4; Indels Score 12.6; DB 1; Pred. No. 4.9e+02; 0; Mismatches 4; 3.0%; Query Match Best Local Similarity 78.9° Matches 15; Conservative

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ò g 575 RESULT

AAS99443 standard; DNA; 20 BP. AAS99443

AAS99443;

12-MAR-2002 (first entry)

cancer related protein 98P7C3 nested PCR primer 2.

Human; 98P6C3; ss; homeodomain protein; vaccine; cytostatic. epitope; transgenic animal; immunogen; T cell; B cell; cytotoxic T cell; CTL; prostate cancer; bladder cancer; kidney cancer; lung cancer; breast cancer; uterine cancer; cervical cancer; stomach cancer; rectal cancer; colon cancer; chromosome 4q11-q12; PCR primer; adapter; suppression subtractive hybridisation; SSH. 

Homo sapiens.

WO200190157-A2.

29-NOV-2001.

24-MAY-2001; 2001WO-US017495.

24-MAY-2000; 2000US-0207138P.

(UROG-) UROGENESYS INC.

Levin E; Challita-Eid PM, Hubert RS, Faris M, Afar DEH, Mitchell SC, Jakobovits A;

WPI; 2002-097642/13.

New isolated 98P7C3-related homeodomain protein highly expressed in various cancers, useful in cancer vaccines and for generating immune response directed to 98P7C3 in mammal.

Example 1; Page 53; 155pp; English.

The invention relates to an isolated 98P7C3-related protein which is a homeodomain protein highly expressed in various cancers. Also include are nomeodomain protein highly expressed in various cancers. Also include are polymuclectides encoding the protein or proteins 078 identical to 98P7C3, a pharmaceutical composition comprising the polymuclectides (including an expression vector comprising the yestor, an anti-98P7C3 antibody, an non-homen transformed with the vector, an anti-98P7C3 antibody, an non-homen transformed with the vector, an anti-98P7C3 antibody, an non-homen transformed with the vector, an anti-98P7C3 antibody, an non-homen transformed with the vector, an anti-98P7C3 antibody of the 98P7C3 protein or polymuclectides in a biological sample, monitoring the presence of cancer in an individual by detecting an elevated level of the 98P7C3 protein or polymuclectides and a pharmaceutical composition comprising a modulator of 98P7C3. 98P7C3 protein, or T call R cell composition mammal) to a 98P7C3 protein, upon contact with a cytotoxic T call (CTL) the immunogen may be a nucleic acid encoding the protein or epitope. The antibody is useful for delivering a cytotoxic agent to a cell that expresses 98P7C3, by conjugating the conjugating the conjugate. The modulator is useful for the antibody-agent conjugate. The modulator is useful for treating a patient with a cancer

that expresses 98P7C3 (e.g. prostate cancer, bladder cancer, kidney cancer, lung cancer, breast cancer, uterine cancer, cervical cancer, eccal cancer and colon cancer), by administering to the patient a vector that comprises the modulator, such that the vector delivers a single chain monoclonal antibody coding sequence to the cancer cells and the encoded single chain antibody is expressed intracellularly in it. The gene for 98P7C3 is located on human chromosome 4q11-q12. The present sequence is oligonucleotide adapter or PCR primer used to isolate a CDNA sequence. hybridisation, SSH

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other; 8888888888888

ó 3.0%; Score 12.6; DB 1; Length 20; 78.9%; Pred. No. 4.9e+02; tive 0; Mismatches 4; Indels 3.0 Best Local Similarity 78.9 Matches 15; Conservative

319 GCGTGCTGGCGGCGACGA 337 2 ecercercececceacea 20 ઠે 셤

RESULT 576

ABK67422 standard; DNA; 20 BP. ABK67422 ID ABK

ABK67422;

(first entry) 02-JUL-2002 Human 83P2H3 cDNA isolation nested PCR primer 2.

Human; human leukocyte antigen; HLA; immunogen; 83P2H3; CaTrF2EH1; calcium transport protein; cancer; prostate cancer; cytostatic; chromosome 7q34; chromosome 12q24.1; T cell; B cell; Bs; primer.

Homo sapiens.

WO200214361-A2.

21-FEB-2002.

17-AUG-2001; 2001WO-US025782.

17-AUG-2000; 2000US-0226329P.

(AGEN-) AGENSYS INC.

, Faris M, Saffran DC, Afar DBH; Jakobovits A; Raitano AB, Challita-Eid PM, Levin E, Hubert RS, Ge W, G

WPI; 2002-269179/31.

Monitoring 83P2H3 gene products for monitoring the presence of cancer i a subject, comprises determining the status of 83P2H3 gene products in tissue sample from the subject and comparing it to a normal sample. 

Example 1; Page 76; 270pp; English.

The invention relates to monitoring 83P2H3 (a calcium transport protein whose gene is located on chromosome 7q34) gene products in a biological sample from an patient who has or is suspected of having cancer cancer; captured of having cancer cancer; captured in a biological sample from an an especially prostate cancer; captured in a tissue sample from an comparing the status to the status of 83P2H3 gene products and (b) comparing the status to the status of 83P2H3 gene products in a normal sample. Also included are modulators of 83P2H3 gene function or status, generating antibodies/immune response against 83P2H3 (or related protein Carrezzil whose gene is located on chromosome carreage protein carreage and elevant sample from an eukocyte antigen) binding peptides derived from the protein, delivering a cytotoxic agent to a cell captured from the protein delivering an antigen-binding region of the antibody, a recombinant protein cantant that produces the recombinant protein, a non-human transgenic animal that produces the recombinant protein, a

tight chains of the anti-appeals the variable domains of the heavy and monoclonal antibody that comprises the variable domains of the heavy and inght chains of the anti-appeals antibody, a vector comprising a polynucleotide that encodes the monoclonal antibody and inducing an immune response to a 8322H3 protein, by providing a 83P2H3-related protein that comprises a T cell or B cell, respectively. The method contacting the protein that comprises a T cell or B cell, respectively. The method is useful for monitoring 83P2H3 gene products in a biological sample for monitoring the growth of cancer in an individual. The modulator is useful for inhibiting the growth of cancer cells that express 83P2H3, for treating an immune response against 83P2H3, and for detecting the cancer that expresses 83P2H3. The immunological methods are useful for generating an immune response against 83P2H3, and for detecting the presence of 83P2H3. The immunological methods are useful for generating an immune response against 83P2H3, and for detecting the presence of 83P2H3. The immunological methodologies and treatment, to detect and quantify 83P2H3 and mutant control of presence of antibodies and treatment, to detect and quantify 83P2H3 and mutant control is sample from a patient mimic the 83P2H3-related protein, for generating anti-italic didotypic antibodies that mimic the 83P2H3 crelated protein. The present sequence is a PCR primer used in the isolation of cDNA encoding 83P2H3 or its 8888888888888888888888888888888

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

ö 0; Gaps Match 3.0%; Score 12.6; DB 1; Length 20; Local Similarity 78.9%; Pred. No. 4.9e+02; les 15; Conservative 0; Mismatches 4; Indels

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ABK70514 standard; DNA; 20 RESULT 577

ABK70514;

S-JUL-2002 (first entry)

Human; cytostatic; 85PlB3; cancer; immunogen; 88; primer; PCR; chromosome 15q14. Human cDNA 85P1B3 nested PCR primer 2.

Homo sapiens.

WO200218578-A2.

07-MAR-2002

28-AUG-2001; 2001WO-US026838.

28-AUG-2000; 2000US-0228432P.

(AGEN-) AGENSYS INC.

Challita-Eid Ge ₩, Faris M, Hubert RS, Afar D, Raitano AB, F Jakobovits A;

WPI; 2002-382963/41.

Composition for modulating the status of 85P1B3 protein or a molecule comprising a substance e.g. antibody specific to, nucleic acid encoding, or ribozyme of 85P1B3.

Example 1; Page 76; 201pp; English.

The invention relates to a composition comprising a substance that modulate the status of 85P1B3, where the status of a cell expresses 85P1B3 gene product is modulated. Also included are a composition 

comprising a peptide region of 5 amino acids of the 85PBB protein, in selected from an az position having a value greater than 0.5 in the selected from an az position having a value greater than 0.5 in the profile, an az position having a value greater than 0.5 in the percent accessible residue profile, an az position having a value greater than 0.5 in the percent accessible residue profile, an az position having a value greater than 0.5 in the percent accessible residue profile, an az position having a value greater than 0.5 in the beta-turn profile; a polymucleotide that encodes analogue peptide of 8, 9, 10 or 11 contiguous cession as protein; a recombinant protein comprising the antigon-binding region of a monoclonal antibody; a non-human transgance antigody (MAD) that binds to the 85PBB protein; a single chain monoclonal antibody (MAD) that comprise the variable domains of the monoclonal antibody (MAD) that encodes the MAD; inhibiting growth of comprising a polymucleotide that encodes the MAD; inhibiting growth of comprising a polymucleotide that encodes the MAD; inhibiting growth of comprision that protein, and generating a patient who bears cancer cells that expresses the protein, and generating a mammalian immune response crobing the protein, and generating a mammalian immune specifically recognize the protein, and generating an ammalian immune specifically recognize the protein, and generating a mammalian immune specifically recognize the protein, and generating a mammalian immune specifically croomyriding a cytoroxic agent to a cell that expresses the protein, conjugated to are independent of the protein, is useful for the protein, is useful for comprises the protein, for detecting the spreases the protein or denominated to antibody and exposing the cell sort treating a patient who bears cancer cells that expresses the protein, for detecting the presence of the protein or expresses the protein, for detecting the presence of the protein or polymucleotide in a biological sample in a patient who has or wh

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

ö Query Match
3.0%; Score 12.6; DB 1; Length 20;
Best Local Similarity 78.9%; Pred. No. 4.9e+02;
Matches 15; Conservative 0; Mismatches 4; Indels

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AAL40496 standard; DNA; 20 BP.

AAL40496;

19-SEP-2002 (first entry)

158P1D7 cDNA related PCR primer SEQ ID No 668.

Cytostatic; 158P1D7; cancer; bladder cancer; mouse; rat; rabbit; dog; cat; cow; horse; human; vaccine; gene therapy; PCR; primer; ss. 

Homo sapiens.

WO200216593-A2.

28-FEB-2002.

22-AUG-2001; 2001WO-US026276

22-AUG-2000; 2000US-0227098P. 10-APR-2001; 2001US-0282739P.

(AGEN-) AGENSYS INC

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Levin
Raitano AB, Afar DEH,
Faris M, Hubert RS, Raitano A
Challita-Eid PM, Jakobovits A;
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12:58:21

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Wed Apr

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WPI; 2002-425659/45.

ö New compositions comprising a gene (designated 158P1D7), its encoded protein or their modulators, useful for treating or diagnosing cancers, particularly bladder cancer, in mammals (e.g. dogs, cats, cows, horses or humans)

Example 1; Page 68; 181pp; English

The invention relates to a novel nucleic acid, designated 158PID7. The compositions are useful for treating or diagnosing cancers, particularly bladder cancer, in mammals (e.g. mice, rats, rabbits, dogs, cats, cows, horses or humans). The compositions are also useful for monitoring genetic abnormalities and in preparing cancer vaccines. The nucleic acid of the invention can be used in gene therapy to treat the said disorders. This polynucleotide sequence represents a PCR primer of the 158PID7 cDNA invention

Seguence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

Gaps ö 3.0%; Score 12.6; DB 1; Length 20; 78.9%; Pred. No. 4.9e+02; tive 0; Mismatches 4; Indels Conservative Similarity 15; Query Match Local Matches

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RESULT 579

AAL53476 standard; DNA; 20 BP AAL53476

(first entry) 16-JAN-2003 AAL53476;

Zinc transporter protein 108P5H8 nested primer 2.

Cytostatic; gene therapy, vaccine, zinc transporter protein 108P5H8; cancer; breast; colon; ovarian; lung; humoral; cellular immune response; passive immunisation; PCR; primer; ss.

Unidentified

WO200260953-A2

17-DEC-2001; 2001WO-US049133.

15-DEC-2000; 2000US-0256210P

(AGEN-) AGENSYS INC.

Challita-Eid PM,

Composition comprising a substance that modulates the status of a zinc transporter protein (108P5H8), useful in diagnosing and treating patients with cancer that express 108P5H8, such as breast, colon, ovarian or lung Hubert RS, Mitchell SC;
Jakobovits A; Eid PM, Faris M, Afar DEH, Morrison KJM, Raitano AB, WPI; 2002-627469/67. Levin E,

Example 1; Page 95; 309pp; English.

The invention relates to a new composition comprising a substance that modulates the status of a zinc transporter protein, designated as 108P5H8, or a molecule that is modulated by 108P5H8. The composition is

useful in diagnosing, preventing, prognosticating or treating patients with cancer that expresses 108P5H8, such as breast, colon, ovarian or lung cancer. The 108P5H8 gene or its fragment can be used to elicit humoral or cellular immune response. The antibodises are useful in active or passive immunisation. The 108P5H8 polymucleotides are useful as probes and primers for the amplification or detection of 108P5H8 genes, as coding sequences for directing the expression of 108P5H8 polypeptides, or as tools for modulating or inhibiting the expression of 108P5H8 genes. The polymucleotides of the invention can be used to treat disorders by gene therapy. This polymucleotide sequence represents a zinc transporter protein 108P5H8 related PCR primer of the invention

88888888888888

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

Gaps ö 3.0%; Score 12.6; DB 1; Length 20; 78.9%; Pred. No. 4.9e+02; ive 0; Mismatches 4; Indels Query Match
Best Local Similarity 78.9
Matches 15; Conservative

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RESULT 580 ABV99876

ABV99876 standard; DNA; 20 BP.

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ABV99876;

(first entry) 28-MAR-2003 Human 121P2A3 post-SSH nested PCR primer 2.

Human; 121P2A3; cytostatic; immunostimulant; vaccine; PCR; primer; humoral immune response; ss; suppression subtractive hybridisation; SSH.

Ношо

WO200283068-A2

24-OCT-2002

10-APR-2001; 2001US-0282739P. 25-APR-2001; 2001US-0286630P. 22-JUN-2001; 2001US-0300373P. 09-APR-2002; 2002WO-US011359. 

(AGEN-) AGENSYS INC

Mitchell SC; W, Jakobovits A; Challita-Bid PM, Raitano AB, Faris M, Hubert RS, Afar DEH, Saffran D, Morrison K, Morrison RK, Ge

WPI; 2003-092956/08.

New composition comprising a substance that modulates the status of 121P2A3 polypeptides, useful for eliciting humoral or cellular immune responses or in assessing the status of 121P2A3 gene products in normal versus cancerous tissues.

Example 1; Page 70; 362pp; English.

The invention relates to a novel composition comprising a substance that modulates the status of a protein, 121P2A3. The composition of the invention has cytostatic and immunostimulant activity, and is useful as a vaccine. The 121P2A3 proteins and polynucleotides are useful for eliciting humoral or cellular immune response. The polynucleotides are useful for characterising cytogenetic abnormalities of this chromosomal locus, as tools that can be used to delineate cytogenetic abnormalities in the chromosomal region that encodes 121P2A3 that may contribute to malignant phenotype, and in assessing the status of 121P2A3 gene products in normal versus cancerous tissues. The proteins are useful for

ABT43860 standard; DNA; 20 BP

RESULT 582

(first entry)

16-OCT-2003

ABT43860;

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generating and characterising domain-specific antibodies, for identifying agents or cellular factors that bind to 121P2A3 or a particular structure domain, and in various therapeutic and diagnostic contexts, including cancer vaccines. The antibodies or I cells reactive with the product are useful in passive or active immunisation, and in imaging methodologies for the management of cancer. The present sequence represents an nested PCR primer used in the invention to amplify gene fragments resulting from suppression subtractive hybridisation (SSH) reactions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention describes a new composition comprising a substance that modulates the status of 158P3D2 or a molecule that is modulated by 158P3D2, where the status of a cell that expresses 158P3D2 is modulated. The composition is useful for treating cancer. This sequence represents novel protein 158P3D2 associated primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New composition comprising a substance that modulates the status of 158P3D2 or a molecule that is modulated by 158P3D2, useful for treating
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                                                                                                                                                                                                     Query Match 3.0%; Score 12.6; DB 1; Length 20; Best Local Similarity 78.9%; Pred. No. 4.9e+02; Matches 15; Conservative 0; Mismatches 4; Indels
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                                                                                                                                                                   Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel protein 158P3D2 associated primer #4.
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                                                                                                                                                                                                                                                                              319 GCGTGCTGGCGGCGGACGA 337
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25-APR-2001; 2001US-0286630P.
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Best Local Similarity 78.9°
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                ACA64671;
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ACA64671
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The invention relates to a novel composition comprising a substance that modulates the status of a 151934 protein (e.g. 151934 variant 1-11; or a molecule that is modulated by the 151934 protein, where the status of a cell that expresses the 151934 protein is modulated. The novel compositions, or the 151934 protein and genes, are useful for eliciting a humoural or cellular immune response. The 151934 genes and proteins are also useful for diagnosing, prognosing preventing or treating cancer, e.g. adenocarcinoma, bladder cancer, colorectal cancer, lung or bronchial cancer, breast cancer or carcinoma. This polynucleotide sequence represents a 1519304 related primer of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New 151P3D4 proteins and genes, useful for eliciting a humoral or cellular immune response, or for diagnosing, prognosing, preventing or treating cancer, e.g. adenocarcinoma, bladder cancer, lung, breast cancer
                                                                                                          Cytostatic; gene therapy; Vaccine; modulator; 151P3D4; humoural; cancer; cellular immune response; adenocarcinoma; bladder; colorectal; lung; bronchlal; breast; carcinoma; PCR; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                      Faris M, Hubert RS, Morrison K;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 69; 426pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                      Challita-Eid PM, Raitano AB, Fari
Morrison RK, Ge W, Jakobovits A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        319 GCGTGCTGGCGGCGACGA 337
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                                                                                                                                                                                                                                                                                           09-APR-2002; 2002WO-US011644.
                                                                        DPNCDN nested primer 2 (NP2).
                                                                                                                                                                                                                                                                                                                               10-APR-2001; 2001US-0282739P.
25-APR-2001; 2001US-0286630P.
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                                                                                                                                                                                                                                                                                                                                                                                    (AGEN-) AGENSYS INC
                                                                                                                                                                                                                    WO200283860-A2.
                                                                                                                                                                                   Unidentified.
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                                                                                                                                                                                                                                                        24-OCT-2002.
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Composition for diagnosing, prognosing, preventing or treating cancer, for eliciting a humoral or cellular immune response, or for active or passive immunization, comprises a substance that modulates the status of a 162P1E6 protein.
transgenic animal; vaccine; gene therapy; PCR; primer; ss.
                                                                                                                                                                                                                      Example 1; Page 71; 437pp; English.
                                                                                                                                        Jakobovits A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              319 GCGTGCTGGCGGCGACGA 337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 GCGTGGTCGCGGCCGAGGA 20
                                                                                                                               Challita-Eid PM, Raitano AB,
Morrison RK, Ge W, Jakobovit
                                                                                      10-APR-2001; 2001US-0283112P.
25-APR-2001; 2001US-0286630P.
                                                                     09-APR-2002; 2002WO-US011544
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                                                                                                               (AGEN-) AGENSYS INC
                                 WO200283916-A2
                 Unidentified
                                                   24-OCT-2002
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Matches
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The invention describes a composition comprising a substance that modulates the status of a protein (1) of 340 or 283 amino acids, or of any of the 15 sequences of 259 amino acids, given in the specification, or a molecule that is modulated by the protein, where the status of the cell that expresses the protein is modulated. The compositions, proteins, polymucleotides and methods are useful for generating and electring cancer. The STEAP-1-related proteins are useful for generating cancer vaccines. The polymucleotides are useful as tools for delineating, with greater precision, cytogenetic abnormalities in the chromosomal region that encodes STEAP-1 that may contribute to the malignant phenotype. This sequence represents a primer used to analyse human six transmembrane epithelial antigen of the prostate or STEAP-1 CDNA's

status of

STEAP-1-related protein, useful for treating and detecting cancer.

Example 1; Page 70; 248pp; English.

New composition comprising a substance that modulates the

WPI; 2003-313240/30.

Faris M, Hubert RS, Morrison K;

Faris M,

(AGEN-) AGENSYS INC

Ge W, Raitano AB, Challita-Eid PM, Jakobovits A;

06-SEP-2002; 2002WO-US028371. 06-SEP-2001; 2001US-0317840P. 05-APR-2002; 2002US-0370387P.

WO2003022995-A2.

20-MAR-2003

Homo sapiens.

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The invention relates to a novel composition comprising a substance that modulates the status of a 162P1E6 protein. The protein comprises one of a 162P1E6 protein. The protein comprises one of a sequences of 70 - 146 amino acids, given in the specification, or a molecule that is modulated by the protein, where the status of a cell that expresses the protein is modulated. An antibody to the 162P1E6 protein is used to deliver a cytotoxic agent or a diagnostic agent to a cell that expresses the 162P1E6 protein. The composition is used to inhibit the growth of cancer cells or generate an immune response. The composition is used for detecting the presence of a 162P1E6 related protein or a 162P1E6-related polymucleotide in a sample. The 162P1E6 proteins and polymucleotides encoding them are useful for diagnosing, proteins and polymucleotides encoding them are useful for diagnosing, prostate cancer, kidney cancer, lung cancer, or breast cancer. They can also be used for eliciting a humoral or cellular immune response. The antibodies or T cells reactive with 162P1E6 are useful for active or passive immunisation. Transgenic animals are useful for developing and screening of useful reagents. The polymucleotide and polympetide sequences of the invention can also be used to treat disorders by being used in a vaccine or in gene therapy. This polymucleotide sequence
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Gaps

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Query Match
3.0%; Score 12.6; DB 1; Length 20;
Best Local Similarity 78.9%; Pred. No. 4.9e+02;
Matches 15; Conservative 0; Mismatches 4; Indels

319 GCGTGCTGGCGGCGGACGA 337

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Seguence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

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Cytostatic; vaccine; cancer; immune response; PCR; primer; ss.
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                                                                                                                                                                                                                               Faris M,
                                                                                                                                                                                                                              Jakobovits A, Challita-Eid PM,
                 ABZ78176 standard; DNA; 20 BP.
                                                                                                                                                           10-APR-2002; 2002WO-US011654.
                                                                                                                                                                           10-APR-2001; 2001US-0282739P.
                                                                                                                                                                                    10-APR-2001; 2001US-0283112P.
25-APR-2001; 2001US-0286630P.
                                                    (first entry)
                                                                                                                                                                                                              (AGEN-) AGENSYS INC.
                                                                     Nested primer #2.
                                                                                                                        WO200283921-A2
                                                    19-MAY-2003
                                                                                                                                        24-OCT-2002.
                                                                                                        Synthetic.
                                   ABZ78176;
RESULT 585
ABZ78176
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STBAP-1; six transmembrane epithelial antigen of the prostate; cancer; cancer vaccine; delineation; cytogenetic abnormality; cytostatic; vaccine; PCR; primer; ss.

Suppressive subtractive hybridisation of STEAP related primer #8.

(first entry)

31-JUL-2003

ACD02621;

ACD02621 standard; DNA; 20 BP.

ACD02621

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Morrison RK, Raitano AB;
    WPI; 2003-075555/07.
Morrison K,
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New composition comprising a substance that modulates the structure of proteins and polynucleotides, useful for therapeutic, prognostic and diagnostic reagents for eliciting cellular or humoral immune response in cancer patients.

Example 1; Page 72; 1021pp; English.

The present invention relates to novel human cancer-related genes and proteins (AB278120-AB278166 and ABR01789-ABR01861). The genes and proteins are useful for eliciting a humoral or cellular immune response. The genes are useful as probes and primers for the amplification and/or detection of genes, mRNAs or their fragments, as reagents for the adjances and/or prognosis of cancer, as coding sequences capable of directing the expression of the protein, as tools for modulating or inhibiting the expression of genes and/or translation of transcripts, and as therapeutic agents. The proteins and peptides are useful as therapeutic, prognostic and diagnostic reagents for cancer. The present sequence is a primer, used in an example from the invention

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

3.0%; Score 12.6; DB 1; Length 20; 78.9%; Pred. No. 4.9e+02; ative 0; Mismatches 4; Indels 319 GCGTGCTGCCGCCGACGA 337 2 GCGTGGTCGCGGCCGAGGA 20 15; Conservative Query Match Best Local Similarity Matches ò ద

ABZ20563 standard; DNA; 20 BP 33-MAR-2003 (first entry) ABZ20563; RESULT 586 ABZ20563 

Cancer associated coding sequence; cancer; human; cytostatic; gene therapy; PCR; primer; 88. Cancer associated coding sequence PCR primer #3.

Homo sapiens.

WO200283920-A2.

24-OCT-2002

10-APR-2002; 2002WO-US011645.

10-APR-2001, 2001US-0282739P. 10-APR-2001; 2001US-02831D2P. 25-APR-2002; 2001US-0286630P. 10-APR-2002; 2002US-00286630.

(AGEN-) AGENSYS INC

WPI; 2003-093030/08.

Jakobovits A, Hubert RS,

Challita-Bid PM;

New pharmaceutical composition for diagnosing, prognosing, preventing or treating cancer, comprises a substance that modulates a nucleic acid sequence, e.g. 105P1B7, 152P1A2B or 156P3A6, or a molecule modulated by the nucleic acid.

Example 1, Page 34; 72pp; English.

The present invention relates to a pharmaceutical composition comprising a substance that modulates the status of a cancer associated nucleic acid sequence such as given in the specification (see ABZ20564-ABZ20575) or a molecule that is modulated by the above nucleic acid sequence, where the status of a cell that expresses the nucleic acid sequence; smodulated. The composition is useful in diagnosing, prognosing, preventing and/or treating cancer. The nucleic acid sequence may be used in monitoring of abourmalisties, in generating and characterising domain-specific antibodies, for identifying agents or cellular factors that bind to a protein, and in therapeutic and diagnostic contexts, such as diagnostic assays, cancer vaccines, and methods of preparing vaccines. The present sequence is a primer used to identify the cancer associated coding sequences suitable to be modulated in the method of the invention

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Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

ö Gaps 4; Indels 0; 3.0%; Score 12.6; DB 1; Length 20; 78.9%; Pred. No. 4.9e+02; Live 0; Mismatches 4; Indels Best Local Similarity 78.9 Matches 15; Conservative Query Match

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184P1E2 gene-specific nested PCR primer #2. AAL52254 standard; DNA; 20 BP (first entry) 16-OCT-2003 AAL52254; 587 

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0; Gaps

Gene therapy, vaccine, 184P1E2, cancer, genetic abnormality, cellular immune response, immunisation, PCR, primer, ss.

WO200283919-A2. Unidentified.

09-APR-2002; 2002WO-US011643. 24-OCT-2002.

10-APR-2001; 2001US-0282739P. 25-APR-2001; 2001US-0286630P.

(AGEN-) AGENSYS INC.

Faris M, Hubert RS, Morrison K; Ge W, Jakobovits A; Raitano AB, Chalitta-Eid PM, Morrison RK,

WPI; 2003-148269/14.

New 184PIE2 polynuclectide encoding a 184PIE2 protein, useful for diagnosing, prognosing, preventing or treating cancer, in eliciting an immune response, and in chromosome mapping.

Example 1; Page 69; 394pp; English.

The invention comprises the amino acid and coding sequence of a 184PIE2 protein. The DNA and protein sequences of the invention are useful for diagnosing, prognosing, preventing and/or treating cancer. The 184PIE2 DNA and protein sequences may also be used to elicit a humoral or a cellular immune response in patients and in monitoring genetic abonormalities. Antibodies raised against the 184PIE2 proteins may be used in active or passive immunisation. The present DNA sequence is used in the exemplification of the invention

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

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121P1F1; 121P1F1 modulation; human; chromosome 4q; cytostatic; gene therapy; vaccine; cancer; immune response; immunisation; primer; ss.

121P1F1 gene nested primer (NP) 2 SEQ ID NO:721.

(first entry)

29-JAN-2004

ADD84533;

ADD84533 standard; DNA; 20 BP.

RESULT 589

ADD8453.

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This invention relates to a novel gene designated 205P1B5, and the encoded protein, which is aberrantly expressed in prostate cancer. Specifically, it refers to the two variants of 205P1B5 mapped to chromosome 821-8912, namely 205P1B4v1 and 205P1B5v2 and fragments thereof that serve as useful diagnostic, prophylactic, prognostic and/ or therapeutic targets for prostate and other types of cancers. The present invention describes methods for the isolation of 205P1B5, for generating an immune response and for generating transgenic or knock out animals for the development and screening of therapeutically useful reagents. Purthermore, it refers to identifying proteins, small molecules or other agents that interact with 205P1B5, and can be used to identify pathways activated by 205P1B5. Accordingly, these are cytostatic and immunogenic compositions that are useful for the development of cancer vaccines. This suppressive subtractive hybridisation (SSH) to isolate the 205P1B5 CDNA fragment of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    205P1B5; prostate cancer; immune response; transgenic; knock out animal; cytostatic; immunogenic; vaccine; ss; SSH; suppressive subtractive hybridisation; PCR; primer; NP2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nested PCR primer 2 (NP2) used in SSH to isolate 205P1B5 cDNA fragment,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New polynucleotide designated 205P1B5, for diagnosing and treating prostate cancer, and as probes or primers for the amplification and/or detection of 205P1B5 genes.
                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Raitano AB, Faris M, Hubert RS, Jakobovits A;
                                                                             ö
3.0%; Score 12.6; DB 1; Length 20; 78.9%; Pred. No. 4.9e+02; ative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 60; 162pp; English.
                                                                                                                                                   319 GCGTGCTGGCGGCGACGA 337
                                                                                                                                                                                                                         2 dceregrededecedada 20
                                                                                                                                                                                                                                                                                                                                                                                                                     ADC71183 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-AUG-2002; 2002WO-US027760.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           31-AUG-2001; 2001US-0316664P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18-DEC-2003 (first entry)
                                    Local Similarity 78.9
Les 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-354484/33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (AGEN-) AGENSYS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Challita-Eid PM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO2003020954-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-MAR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADC71183;
        Query Match
                                                                                                                                                                                                                                                                                                                                       RESULT 588
                                                                             Matches
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Composition comprising a substance that modulates the status of 121P1F1, useful in diagnosing, preventing, prognosticating or treating patients with cancer that expresses 121P1F1, such as breast, colon, ovarian or

Ge ⊠;

Challita-Eid PM, Hubert RS, Raitano AB, Faris M, Afar DEH,

WPI; 2003-156757/15.

lung cancer.

Jakobovits A;

28-FEB-2002; 2002WO-US006242. 05-MAR-2001; 2001US-00799250

WO200295009-A2.

28-NOV-2002.

Homo sapiens.

Synthetic

(AGEN-) AGENSYS INC

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The present invention describes a composition (I) comprising a substance that modulates the status of 121PIFI (gene and encoded protein), or a molecule that is modulated by 121PIFI (gene and encoded protein), or a molecule that is modulated. The human 121PIFI gene maps to chromosome 4q. (I) has cytostatic activity, and can be used in gene therapy, and in vacines. The composition (I) can be used for diagnosing, preventing, prognosticating or treating patients with cancer that expresses 121PIFI, such as breast, colon, ovarian or lung cancer. The 121PIFI gene or its cragment can be used to alicit a humoral or cellular immune response. If IPIFI antibodies can be used in active or passive immunisation. 121PIFI polynoclecides are useful as probes and primers for the amplification or detection of 121PIFI pelpeptides, or as coding sequences for the amplification or inhibiting the expression of 121PIFI genes. The present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.0%; Score 12.6; DB 1; Length 20; 78.9%; Pred. No. 4.9e+02; tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Seguence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Page 71; 285pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       319 GCGTGCTGGCGGCGGACGA 337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2 ecercercececeacea 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADE65924 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-JAN-2004
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Gaps

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Query Match 3.0%; Score 12.6; DB 1; Length 20; Best Local Similarity 78.9%; Pred. No. 4.9e+02; Matches 15; Conservative 0; Mismatches 4; Indels

319 GCGTGCTGGCGGCGGACGA 337

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2 sceregrescesceases 20

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

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Query Match 3.0%; Score 12.6; DB 1; Length 20; Best Local Similarity 78.9%; Pred. No. 4.9e+02; Matches 15; Conservative 0; Mismatches 4; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                      This invention relates to a novel composition which comprises a substance that modulates the status of a novel protein (16122108) and its variants having a sequence of 875 amino acids provided in the specification. The protein of the invention is over-expressed in certain cancers. The compounds of the invention may have cytostatic activity and the sequence of the lelpzing protein, and the gene which encodes it, may be useful for gene therapy or the development of a vaccine. The composition and methods of the invention are useful in diagnosing, preventing and retacting acucer. The present sequence is that of PCR primer which was used for amplification of a region of the gene encoding the human lelpziloB protein during the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                            A composition for diagnosing, preventing and treating cancer (e.g. prostatic, renal or uterine cancer) comprises 161P2F10B polynucleotides and polypeptides.
                                                                                                                                                                                                                                                          Jakobovits A, Raitano AB, Faris M, Hubert RS, Ge W, Morrison KJM;
Morrison RK, Challita-Eid PM;
                             161P2F10B; cancer; cytostatic; gene therapy; vaccine; PCR; primer; ss;
Human 161P2F10B protein-related PCR primer SegID36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                               Example 1; SEQ ID NO 36; 135pp; English
                                                                                                                                                                                      07-NOV-2001; 2001US-00005480.
                                                                                                                                                             07-NOV-2002; 2002WO-US036002
                                                                                                                                                                                                                                                                                                           WPI; 2003-441560/41.
                                                                                                                                                                                                                                  (AGEN-) AGENSYS INC.
                                                                                                   WO2003040340-A2
                                                                         Homo sapiens.
                                                                                                                               15-MAY-2003
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This invention relates to novel composition comprising a substance that modulates the status of a 433 residue protein, given in the specification with the DNA sequence encoding it, or a molecule that is modulated by the protein. The novel protein 193PEIB exhibits tissue specific expression in normal adult tissue and is aberrantly expressed in cartain cancers. Compositions which modulate the 193PIBIB protein may have cytostatic activity and the DNA sequence which encodes protein 193PIBIB may be useful in gene therapy. The composition of the invention may be useful por the treatment of cancer. The present sequence is that of an RT-PCR primer which was used for the amplification of human 193PIBIB gene DNA during the exemplification of the invention.

for

3 ę,

Hubert RS,

Faris M,

Challita-Eid PM,

Raitano AB, Jakobovits A

WPI; 2003-532905/50.

07-DEC-2001; 2001US-00013312

(AGEN-) AGENSYS INC

New composition comprising 193PIE18-related protein, useful preventing or treating cancer.

Example 1; SEQ ID NO 59; 260pp; English

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Gaps

4; Indels 0;

Seguence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

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Typing nucleic acid for obtaining information about several variable sites involves simultaneously or sequentially performing two or more primer extension reactions, and determining the pattern of inucleotide
                                                                                                                                                 Single nucleotide polymorphism; nucleic acid typing; tissue typing; human; PCR; primer; angiotensin converting enzyme; ACE; ss.
                                                                                                                              Angiotensin converting enzyme SNP fragment Bu6 PCR primer B.
                                                                                                                                                                                                                                                                                                                         Pourmand N;
                                                                                                                                                                                                                                                                                (STRD ) UNIV LELAND STANFORD JUNIOR. (GARD/) GARDNER R.
319 GCGTGCTGGCGCGGACGA 337
                                              2 GCGTGGTCGCGGCCGAGGA 20
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193P1E1B; tissue specific expression; cancer; cytostatic; gene therapy; cancer; human; PCR; RT-PCR; reverse transcription PCR; primer; ss.

06-DEC-2002; 2002WO-US039274

WO2003050255-A2

19-JUN-2003

Homo sapiens,

Juman protein 193P1E1B-related PCR primer SeqID59.

(first entry)

29-JAN-2004

ADD96944;

ВР

ADD96944 standard; DNA; 20

RESULT 591

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% Match 3.0%; Score 12.6; DB 1; Length 20; Local Similarity 78.9%; Pred. No. 4.9e+02; les 15; Conservative 0; Mismatches 4; Indels

Query Match Best Loca Matches

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Page 285
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The invention relates to a novel method for obtaining typing information about several variable sites within target nucleic acid, or typing one or mote nucleic acid molecules. The methods of the invention are useful for typing one or more nucleic acid molecules containing two or more variable sites, preferably nucleic acid molecules containing three or more variable sites are typed, where three or more primer exension reactions or performed. The method is also useful for diagnosis of pathological conditions characterized by the presence of specific nucleic acid molecule(s). The methods are particularly suited for identifying microbial species or their subtypes, and in typing procedures e.g. typing of polymorphisms, tissue typing or in clinical applications. The sequence connected primer used in the invention to amplify the single nucleotide polymorphism (SNP) Eu6 from the angiotensin converting enzyme (ACE) gene. The primer binds to the template with its 3, end 5 cucleotides from the SNP position
                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Enzymatic nucleic acid; ribozyme; trans cleavage; inhibition; gene expression; downregulation; interleukin-5; IL-5; ICAM-1; intercellular adhesion molecule; rel A; tumour necrosis factor; TNF-alpha: respiratory syncytial virus; RSV; bcr-ab; oncogene; translocation; chronic myelogenous leukaemia; CML; cancer; Philadelphia chromosome; inflammation; autoimmune disease; atherosclerosis; myocafdial infarction; stroke; restenosis; transplant rejection; rheumatoid arthritis; psoriasis; myocafdial; Kawasaki disease; septic shock; HIV; human immunodeficiency virus; acquired immune deficiency syndrome; AIDS;
                                                                                                                                                                                                                                                                                                                                                                                                      0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human relA hammerhead ribozyme target sequence (nt. position 1058).
                                                                                                                                                                                                                                                                                                                                                                 Query Match 2.9%; Score 12.4; DB 1; Length 14; Best Local Similarity 92.9%; Pred. No. 2.5e+02; Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                              Sequence 14 BP; 1 A; 5 C; 4 G; 4 T; 0 U; 0 Other;
Disclosure; Fig 5B; 86pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     94US-00201109.
94US-00218934.
94US-00227488.
94US-00227958.
94US-00227958.
94US-00245736.
94US-0021932.
94US-00291433.
94US-00292620.
94US-00292620.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5127/c
AAT55127 standard; RNA; 15 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                         268 ACCTGGAGCAGGGC 281
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-MAR-2003 (revised)
21-APR-1997 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14 ACCTGGAGCAGAGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
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29-MAR-1994;
04-APR-1994;
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19-AUG-1994;
02-SEP-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAT55127;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 593
AAT55127/c
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The present sequence represents a preferred target sequence for an enzymatic nucleic acid (i.e. a ribozyme) which cleaves relA mRNA at the nucleotide base position indicated in the DE line. The relA gene product is a subunit of the transcriptional regulator NF-kappaB and is implicated specifically in the induction of inflammatory responses. Regions of the mRNA that do not form secondary folding structures and that contain protential hammerhead and hairpin ribozyme cleavage sites were identified by computer analysis. Ribozymes directed against these mRNA sequences were designed and synthesised with modifications that improve their nuclease resistance. The ribozymes are designed to cleave the target sequences and thereby inhibit relA expression, making them potentially useful for treating rheumatoid arthritis, restenosis and asthma as well as for increasing tolerance to transplanted tissues. The potential in immunosuppressive properties of a ribozyme that cleaves relA mRNA means that uses are limited to local delivery, acute indications or ex vivo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         o DT, Chowrira B, Direnzo A, Draper KG, Dudycz LW;
Karpelsky A, Kisich K, Matulic-Adamic J, Mcswiggen JA;
Pavco P, Belgleman L, Sullivan SM, Sweedler D, Thompson JD;
Usman N, Wincott FE, Woolf T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ribozymes having modified bases and methods for producing them - for use in inhibiting disease related genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 2.9%; Score 12.4; DB 1; Length 15; Best Local Similarity 92.9%; Pred. No. 2.9e+02; Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human B7-1 hammerhead ribozyme target SEQ ID NO:1186.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 2; Page 229; 407pp; English.
94US-00303039.
94US-00311486.
94US-00311749.
94US-00316771.
94US-00334847.
94US-00334847.
94US-00334847.
94US-0033593.
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                                                                                                                                                                                                                                                                                                                                                                      94US-00363233
                                                                                                                                                                                                                                                                                                                                                                                                  95US-00380734
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-JUL-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14 GAGGTGGAGGCCGG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Stinchcomb DT,
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                                                                                              28-SEP-1994;
03-OCT-1994;
07-OCT-1994;
04-NOV-1994;
10-NOV-1994;
16-DRC-1994;
23-DEC-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Grimm S, K
Modak A, F
Tracz D, U
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       $\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac
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neutral lipid transfer; plasma lipoprotein; atherosclerosis; athereccomy; reverse cholesterol transport; high density lipoprotein; therapy; CETP; familial hypercholesterolosemia; dyslipideamia; hypoalphalipoproteinaemia; peripheral vascular disease; hyperbetalipoproteinaemia; RCT; inhibitor; angioplastic restenosis; low density lipoprotein; diabetes; HDL; human;

Hammerhead ribozyme; cholesterol ester transfer protein; mRNA cleavage;

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The present invention describes a novel enzymatic nucleic acid (ENA) having a hammerhead motif (HM) comprising; (i) at least 5 ribose residues 7 (ii) a 2-C-allyl modification at position 4 of the ENA, (iii) at least to 7 (iii) at least 5 ribose residues 5 (iii) at least 5 ribose residues 6 (iv) a 3'-end modification. The ENA's can inhibit collagenase and stromelysin production in the synovial membrane of joints for the treatment or prevention of arthritis, particularly osteoarthritis or rheumatoid arthritis. The ENA's can also be used to treat antigen presenting cells of a donor to induce tolerance in a recipient to an alloantigen of a donor. They can also be used for enhancing graft tolerance or for treating autoimmune disease, and for treating allergies and other inflammatory conditions. The ENA's can also be used in diagnosis. Ribozyme therapy impacts on the expression of a tromelysin without introducing the non-specific effects upon gene expression which accompany treatment with retinoids and dexamethasone. The concentration of ribozyme required to affect a therapeutic treatment is lower than that required of antisense molecules, and is highly present in the present sequence is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Enzymatic nucleic acid molecules having a hammer-head motif - used for
the treatment of arthritis, induction of graft tolerance or treatment of
auto-immune diseases.
                                                                                                                                                                                                                                                                                                                                                                    f, Draper K, Pavco P;
Wincott F, Matulic-Adamic J;
Burgin A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 12.4; DB 1; Length 15;
Pred. No. 2.9e+02;
4; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 15 BP; 2 A; 3 C; 6 G; 0 T; 4 U; 0 Other
                                                                                                                                                                                                                                                                                                                                                                        Jarvis T,
                                                                                                                                                                                                                                                                                                                                                                                       Usman N,
Modak A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 10; Page 166; 307pp; English.
                                                                                                                                                                                                                                                                                                                                                                    Stinchcomb DT,
Gustofson J, Us
Thompson JD, Mc
                                                                                                                    94US-00354920.
94US-00363253.
94US-00363254.
95US-00426124.
95US-00434509.
95US-00434509.
                                                                                        95WO-US015516
                                                                                                                                                                                                                                                                                 95US-00512861
95US-00541365
                                                                                                                                                                                                                                                                                                                                    (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1996-300653/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         present invention
                                                                                                                                                                                                                                                                                                                                                                    Beigelman L,
Mcswiggen J,
Karpeisky A,
                                                                                                                                                                                         20-APR-1995;
02-MAY-1995;
04-MAY-1995;
                    WO9618736-A2
                                                                                      22-NOV-1995;
                                                                                                                                                                                                                                                                                                  05-OCT-1995
                                                                                                                                                       23-DEC-1994
17-FEB-1995
                                                    20-JUN-1996
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New ribozyme(s) for cleaving cholesterol ester transfer protein mRNA -useful for preventing or treating initial development, progression or regression of vascular diseases, esp. familial hypercholesterolaemia.

Claim 4; Page 30; 72pp; English.

Couture L, Stinchcomb D, Mcswiggen J, Bisgaler C,

WPI; 1996-321852/32.

94US-00363240. 95WO-US016000.

11-DEC-1995; 23-DEC-1994;

04-JUL-1996.

Homo sapiens WO9620279-A1

LDL; ss.

(RIBO-) RIBOZYME PHARM INC (WARN ) WARNER LAMBERT CO.

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AAT49608-T49863 represent target sequences for the human cholesterol ester transfer protein (GETP) hammerhead (HH) ribozymes (see AAT49881-CT50137). GETP is a 74 kD glycoprotein that facilitates neutral lipid transfer between plasma lipoproteins. The numbering of the targets refers to the position of the cleavage site in full length GETP. The ribozyme Dinds to Simmediately upstream ther side of this site, provided the sequence UH is immediately upstream. The ribozymes are able to cleave mRNA from the gene encoding GETP, thereby blocking synthesis and/or expression of the mRNA. By inhibiting CETP, the reverse cholesterol transport (RCT) pathway can be inhibited (or eliminated) thereby HDL), prolonging HDL half life, and therefore increasing HDL levels. The ribozymes can be used to treat conditions associated with abnormal levels of CETP, specifically familial hyperbelaipoproteinaemia, atherosclerosis, peripheral vascular disease, hyperbelaipoproteinaemia, hypozaphalipoproteinaemia, dyslipidaemia, associated with abnormal levels of CETP, specifically familial hyperbelaipoproteinaemia, hypozaphalipoproteinaemia, dyslipidaemia, castenosis. Sy inhibiting CETP, the levels of HDL and low angioplastic restenosis. By inhibiting CETP, the levels of HDL and low density lipoproteins (LDL), and the HDL:LDL ratio are favourably altered consity libozofens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (a decrease in LDL levels, and a corresponding increase in HDL levels). The HH ribozymes can also be used diagnostically to study genetic drift and mutations in diseased cells, and to detect CETP mRNA. As the HH ribozymes target specific regions of the CETP gene, they have low non-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 2.9%; Score 12.4; DB 1; Length 15; Best Local Similarity 85.7%; Pred. No. 2.9e+02; Matches 12; Conservative 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 15 BP; 5 A; 3 C; 5 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   엺
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAT49707 standard; RNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Accademocaaceca 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            specific activity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAT49707;
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ID AAT4
XX AAT4
AC AAT4
XX DT 02-M
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Gaps

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h 2.9%; Similarity 64.3%; 9; Conservative '

Query Match Best Local Similarity Matches 9; Conserv

398 GAAGGICTICTACG 411

|| ||:|::|:||| GAGGGUCUUCUACG 15

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Human CETP HH ribozyme target sequence #1056.

(first entry)

02-MAR-1997

AAT49705;

AAT49705 ID AAT4 XX AAT4 AC AAT4 XX DT 02-M XX XX

AAT49705 standard; RNA; 15 BP

RESULT 595

(first entry)

AAV66430;

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Hammerhead ribozyme; cholesterol ester transfer protein; mRNA cleavage; neutral lipid transfer; plasma lipoprotein; atherosolerosis; atherectomy; reverse cholesterol transport; high density lipoprotein; therapy; CETP; familial hypercholesterolaemia; dyslipidaemia; hypoalphalipoproteinaemia; peripheral vascular disease; hyperbetalipoproteinaemia; RCT; inhibitor; angioplastic restenosis; low density lipoprotein; diabetes; HDL; human;
                                                                                                                                                                                                                                                                                                                        New ribozyme(s) for cleaving cholesterol ester transfer protein mRNA useful for preventing or treating initial development, progression or regression of vascular diseases, esp. familial hypercholesterolaemia.
                                                                                                                                                                                                                                                                               Couture L, Stinchcomb D, Mcswiggen J, Bisgaier C,
         Human CETP HH ribozyme target sequence #1057.
                                                                                                                                                                                                                                                                                                                                                                          Claim 4; Page 30; 72pp; English.
                                                                                                                                                                                                                        94US-00363240.
                                                                                                                                                                                                 95WO-US016000.
                                                                                                                                                                                                                                            (RIBO-) RIBOZYME PHARM INC (WARN ) WARNER LAMBERT CO.
                                                                                                                                                                                                                                                                                                       WPI; 1996-321852/32.
                                                                                                                             Homo sapiens
                                                                                                                                                    WO9620279-A1
                                                                                                                                                                                                 .1-DEC-1995;
                                                                                                                                                                                                                        3-DEC-1994;
                                                                                                                                                                         04-JUL-1996.
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Pape M;

AAT 49608-T49863 represent target sequences for the human cholesterol ester transfer protein (CETP) hammerhead (HH) ribozymes (see AAT49881-510137). CETP: is a 74 kD glycoprotein that facilitates neutral lipid transfer between plasma lipoproteins. The numbering of the targets refers to the position of the cleavage site in full length CETP. The ribozyme binds to 5 nucleotides either side of this site, provided the sequence UH is immediately upstream. The ribozymes are able to cleave mRNA from the gene encoding CETP, thereby blocking synthesis and/or expression of the sequence of mRNA. By inhibiting CETP, the reverse cholesterol transport (RCT) pathway can be inhibited (or eliminated) thereby preventing the reduction in size density of the high density lipoproteins (HDL), prolonging HDL half life, and therefore increasing HDL levels of CETP, specifically familial conditions associated with abnormal levels of CETP, specifically familial hypercholesterolaemia, atherosolerosis, peripheral vascular disease, hyperbetalipoproteins (HDL), and the HDL:LDL ratio are favourably altered angioplastic restenosis. By inhibiting CETP, the levels of HDL and low density lipoproteins (LDL), and the HDL:LDL ratio are favourably altered and mutations in diseased cells, and to detect CETP mRNA, As the HH contains in diseased cells, and to detect CETP mRNA, As the HH contains in diseased cells, and to detect CETP mRNA, as the HH contains in diseased cells, and to detect CETP mRNA, as the HH contains in diseased cells, and to detect CETP mRNA, as the HH contains in diseased cells, and to detect CETP mRNA, as the HH contains in diseased cells, and the CETP gene, they have low nonspecific activity

Seguence 15 BP; 5 A; 3 C; 5 G; 0 T; 2 U; 0 Other;

Gaps ö / Match 2.9%; Score 12.4; DB 1; Length 15; Local Similarity 85.7%; Pred. No. 2.9e+02; nes 12; Conservative 1; Mismatches 1; Indels Query Match Matches

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AAV66430 standard; DNA; 15 RESULT 597 AAV66430 ID AAV6 XX

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functioning as promoters for the tetracycline resistance (TetR) gene.

functioning as promoters for the tetracycline resistance (TetR) gene.

They are derived from the -35 promoter sequence of the TetR gene of

They are derived from the -35 promoter sequence of the TetR gene of

The specification describes a method for obtaining an oligonucleotide

that confers a predetermined biological function, such as regulation of

expression or a biological activity of a polypeptide, on a cell. The

method comprises cloning a heterogeneous pool of oligonucleotides into an

expression vector, where the clones oligonucleotides are transcribed or

act as regulatory sequences, introducing a random sample of the cloned

coligonucleotides into a population of cells that do not exhibit the

predetermined biological function, selecting a subpopulation of cells

exhibiting the predetermined biological function, and isolating an

coligonucleotide that confers this function from the selected

subpopulation of cells. The process is used, for example, for identifying

new forms of the Escherichia coli tetracycline resistance gene promoter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Identification of biologically active DNA sequences - by transforming cells with random oligo-nucleotide (s).
                                                                                                        -35 promoter; plasmid pBR322; tetracycline resistance gene; TetR; promoter; Escherichia coli; active site; beta-lactamase gene; ss.
                                                                      -35 promoter sequence of TetR gene of plasmid pBA6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 15 BP; 0 A; 5 C; 9 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    the active site of the beta-lactamase gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Fig 3; 24pp; English.
                                                                                                                                                                                                                                                                                                                                         89US-00368674.
92US-00881607.
93US-00105108.
                                                                                                                                                                                                                                                                                                                         86US-00887070
                                                                                                                                                                                                                                                                                   94US-00316415
                                                                                                                                                                                                                                                                                                                                                                                                                    (UNIW ) UNIV WASHINGTON.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Loeb LA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1998-582545/49.
                                                                                                                                                                                                                                                                                                                     17-JUL-1986;
19-JUN-1989;
12-MAY-1992;
                                                                                                                                                                                                                                                                                   30-SEP-1994;
                                        06-JAN-1999
                                                                            Substituted
                                                                                                                                                                                                                                                                                                                                                                                    11-AUG-1993;
                                                                                                                                                                                                           US5824469-A.
                                                                                                                                                                                                                                                 20-OCT-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Horwitz MS,
                                                                                                                                                                      Synthetic
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.. 0 2.9%; Score 12.4; DB 1; Length 15; 92.9%; Pred. No. 2.9e+02; iive 0; Mismatches 1; Indel8 ВЪ. 143 GGCGGTGGAGGCCG 156 AAC73241 standard; DNA; 15 1 GGCGGTGGCGCCG 14 Best Local Similarity 92.9 Matches 13; Conservative 02-FEB-2001 AAC73241; Query Match RESULT 598 2×2×2×2×2×2 ઠ 셤

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Gape

Oligonucleotide array; genotyping; single base extension reaction; SBE; Forward primer #43 used in multiplexing PCR/SBE assay.

21

Wed Apr

99US-0140345P

21-JUN-1999;

21-JUN-2000; 2000WO-AU000693

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The present invention relates to an oligonucleotide array comprising oligonucleotide tags fixed to a solid substrate. The oligonucleotide array is useful for genotyping a nucleic acid sample at one or more locivia single base extension (SBE) reactions. A pair of primers is used to amplify a polymorphic locus in a sample e.g. a single nucleotide polymorphism (SNP). The present sequence is one of the primers used in the method of the present invention to amplify a polymorphic sample. The amplified nucleic acid product is then used as a template in a SBE reaction with an extension primer. The SBE reaction products are used to form the oligonucleotide array
                                                                                                                                                                                                                                                                                                                                                                                   Universal array of oligonucleotides tags attached to a solid substrate along with locus-specific tagged oligonucleotides useful in genotyping using single base extension reactions.
                                                                                                                                                                                                                                                                                             Hirschhorn JN, Huang X, Kaplan P, Lander ES, Lockhart DJ;
PCR primer; polymorphic locus; single nucleotide polymorphism; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 15 BP; 4 A; 5 C; 5 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                      (WHED ) WHITEHEAD INST BIOMEDICAL RES (AFFY-) AFFYMETRIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 7; Page 52; 70pp; English
                                                                                                                                                                                 99US-0126473P.
99US-0140359P.
                                                                                                                                              27-MAR-2000; 2000WO-US008069
                                                                                                                                                                                                                                                                                                                                               WPI; 2000-656171/63.
                                                                                                                                                                                                                                                                                             Fan J, Hirschhorn
Ryder T, Sklar P;
                                                                        WO200058516-A2
                                                                                                                                                                                     26-MAR-1999;
                                                                                                                                                                                                     23 -JUN-1999;
                                       Unidentified
                                                                                                             05-OCT-2000
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Gaps
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o
Score 12.4; DB 1; Length 15; Pred. No. 2.9e+02; 0; Mismatches 1; Indels
   Query Match
Best Local Similarity 92.9%;
Matches 13; Conservative
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388 ACGGCCCAAGAAG 401 1 ACGGCGCCAAGATG 14 ð 셤

AAF49243 standard; DNA; 15 BP IGF-I oligonucleotide #203 (first entry) 30-MAR-2001 AAF49243; RESULT 599 AAF49243 

Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin discorder; insulin-like Growth Factor. I receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; kearcosis; neophasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplamia; kidney disease; neobascular condition; hyperplamia; kidney disease;

Homo sapiens

WO200078341-A1

28-DEC-2000

The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonucleoride, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, conjugance and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotide which can be used to design the antisense oligonucleotide which can be useful for ameliorating the effects of psoriasis, chthyosis, pityriasis, ubb, pliaris, serborrhoea, keloids, keratosis, ichthyosis, pityriasis, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the skin, a chyperneovascular design and proper of the inside of blood sease, phyperproliferation of the inside of blood seases and the property of the inside of blood seases and the property of the inside of blood seases. Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or Sequence 15 BP; 2 A; 3 C; 7 G; 3 T; 0 U; 0 Other; Edmondson SR; (MURD-) MURDOCH CHILDRENS RES INST. Example 8; Page 62; 201pp; English vessels or any other hyperplasia Werther GA, WPI; 2001-041421/05 inflammation. Wraight CJ, 

Gaps ö Query Match

2.9%; Score 12.4; DB 1; Length 15;
Best Local Similarity 92.9%; Pred. No. 2.9e+02;
Matches 13; Conservative 0; Mismatches 1; Indels

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RESULT 600

AAF53588 standard; DNA; 15 BP. IGF-I oligonucleotide #4548. (first entry) 30-MAR-2001 AAF53588;

Antisense therapy, antiproliferative, antinflammatory, antipsoriatic, cytostatic, dermatological, cardiant, virucide, ophthalmological, keloid, skin discorder, Insulin-like Growth Factor I receptor, IGF-1, pityriasis, IGF binding protein, IGFB-2, IGFBP3, inflammation, psoriasis, plazis, growth factor mediated cell proliferation, ichthyosis, serborrhoea, ruba, keratosis, neoplasia, scleroderma, wart, skin cancer, sclerotic disease, hyperneovascular condition, hyperplasia, kidney disease, neovascular condition, hyperplasia, kidney disease,

21-JUN-2000; 2000WO-AU000693. 99US-0140345P. MO200078341-A1. 21-JUN-1999; Homo sapiens 28-DEC-2000. 

The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligomucleotide, (for Insulin-like Growth Factor [167]-1 receptor, 1GF binding protein [1678]-2 or 1GFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inhibiting or reducing growth factor mediated cell proliferation, configuration and/or other disorders. The present sequence is an oligomucleotide which can be used to design the antisense oligomucleotide which can be useful for ameliorating the effects of psoriasis, season playsis, plyriasis, ruba, pilaris, senborrhoea, keloids, keratosis, neoplasis, soleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, chain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood to sessels or any other hyperplasia Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or Werther GA, Edmondson SR; Example 8; Page 90; 201pp; English. (MURD-) MURDOCH CHILDRENS RES INST WPI; 2001-041421/05. inflammation. Wraight CJ, 

Query Match 2.9%; Score 12.4; DB 1; Length 15; Best Local Similarity 92.9%; Pred. No. 2.9e+02; Matches 13; Conservative 0; Mismatches 1; Indele 356 CAGCGACTTCCTCA 369 15 CAGCCACTTCCTCA 2 ઠે 윱

Sequence 15 BP; 3 A; 1 C; 7 G; 4 T; 0 U; 0 Other;

0; Gaps

AAF53590 standard; DNA; 15 BP. IGF-I oligonucleotide #4550. 30-MAR-2001 (first entry) AAF53590; RESULT 601 AAF53590, 

Antisense therapy; antiproliferative; antinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding procein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition; hyperplasia; kidney disease;

Homo sapiens.

WO200078341-A1 28-DEC-2000 21-JUN-2000; 2000WO-AUD00693.

99US-0140345P. 21-JUN-1999; (MURD-) MURDOCH CHILDRENS RES INST.

Werther GA, Edmondson SR; Wraight CJ,

WPI; 2001-041421/05.

Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation.

Example 8; Page 90; 201pp; English.

The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an anticontection (for Insulin-like Growth Factor IIGF]—I receptor, IGF binding protein [IGF8]—2 or IGF8P3), which is capable of inhibiting or reducing growth factor mediated cell proliferation.

Inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the artisense oligonucleotide which can be used to design the artisense oligonucleotide which can be used to design the artisense oligonucleotide is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperpoliferation of the inside of blood

Sequence 15 BP; 3 A; 2 C; 7 G; 3 T; 0 U; 0 Other;

0; Gaps 2.9%; Score 12.4; DB 1; Length 15; 92.9%; Pred. No. 2.9e+02; ative 0; Mismatches 1; Indels 2.9% Best Local Similarity 92.9% Matches 13, Conservative

355 ACAGCGACTTCCTC 368 14 Acadecaerreere 1 ò

AAF49244 standard; DNA; 15 BP. RESULT 602 AAF49244

AAF49244;

IGF-I oligonucleotide #204. (first entry) 30-MAR-2001 

Antiense therapy, antiproliferative, antinflammatory, antipeoriatic, cytostatic, dermatological, cardiant; virucide, ophthalmological, keloid, skin discorder, ineulin-like Growth Pactor. I receptor; IGF-1; pityriasis; IGF binding proctein, IGFB-2; IGFBP3; inflammation, psoriasis; pitowth factor mediated cell proliferation; ichthyosis; serbornhoea; ruba, keratosis, neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperpleasis; kidney disease; neovascular condition of the retina; ss.

Ношо варіепв.

WO200078341-A1.

28-DEC-2000.

21-JUN-2000; 2000WO-AU000693

99US-0140345P. 21-JUN-1999; (MURD-) MURDOCH CHILDRENS RES INST.

Edmondson SR, Werther GA, Wraight CJ,

WPI; 2001-041421/05.

Ameliorating the effects of a disorder, e.g. psoriasis, by administering

UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation.

Example 8; Page 62; 201pp; English.

The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligomucleotide, (for Insulan-like Growth Factor [IGF].

receptor, IGF binding protein [IGFBP] 2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inhibiting or reducing growth factor mediated cell proliferation, oligomucleotide which can be used to design the antisense oligomucleotides of the present invention (see AAF4151 and AAF45153-F45161). The method is useful for ameliotating the effects of psoriasis, ichthyosis, pityriasis, ruba, pliaris, serborrhoea, keloids, keratosis, ineoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood vessels or any other hyperplasia

Sequence 15 BP; 3 A; 2 C; 7 G; 3 T; 0 U; 0 Other;

0; Gaps Ouery Match 2.9%; Score 12.4; DB 1; Length 15; Best Local Similarity 92.9%; Pred. No. 2.9e+02; Matches 13; Conservative 0; Mismatches 1; Indels

20 GGTGACCGAGGGCT 33

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RESULT 603 AAF49333

AAF49333 standard; DNA; 15 BP. AAF49333; 

IGF-I oligonucleotide #293.

30-MAR-2001 (first entry)

Antisense therapy, antiproliferative, antinflammatory, antipsoriatic, cytostatic, dermatological, cardiant; virucide, ophthalmological, keloid, skin disorder, Insulin-like Growth Factor I receptor; IGF-1; pityriasis, IGF binding procein, IGFBP-2; IGFBP3, inflammation, psoriasis, pracein, IGFBP-2; IGFBP3, inflammation, psoriasis, planis; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neophasia; sclaroderma; wart; skin cancer; selerotic disease; hypermeowascular condition; hyperplasis; kidney disease; neovascular condition of the retina; ss.

Homo sapiens.

WO200078341-A1.

28-DEC-2000.

21-JUN-2000; 2000WO-AU000693.

(MURD-) MURDOCH CHILDRENS RES INST. 99US-0140345P, 21-JUN-1999;

Wraight CJ, Werther GA, Edmondson SR;

WPI; 2001-041421/05.

Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation.

plassing plassing plassing second and or skin, growth increase, hyperplassing isease, kidney disease, hyperplassing sequence 15 BP; 2 A; 3 C; 4 G; 6 T; 0 U; 0 Other;

Auery Match
Best Local Similarity 92.9%; Pred, No. 2.9e+02;
Matches 13; Conservative 0; Mismatches 1; Indels 0; Gaps 0 The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises concacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see AAF45151 and AAF45153-F45161). The method is useful for ameliocating the effects of psoriasis, inhyposis, pityriasis, blairs, serborrhoea, keloids, keratosis, inhyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperpriorieferation of the inside of blood Sequence 15 BP; 2 A; 3 C; 4 G; 6 T; 0 U; 0 Other; Example 8; Page 62; 201pp; English. %\$5555555555555555**%**%

Query Match Best Loca Matches

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AAF49334 standard; DNA; 15 BP. AAF49334; RESULT 604 AAF49334/c

30-MAR-2001 (first entry) 

IGF-I oligonucleotide #294.

Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; insulin-like Growth Pactor I receptor; IGP-1; pityriasis; IGF binding protein; IGFBP-3; inflammation; psoriasis; pityriasis; growth factor mediated cell proliferation; ichthyosis; serborrhoes; ruba; keratosis, neclasis, necladeria; salaroderia; wart; skin cancer; selerotic disease; hyperneovascular condition; hyperplasis; kidney disease; neovascular condition of the retina; ss.

WO200078341-A1. 28-DEC-2000. 21-JUN-2000; 2000WO-AU000693.

99US-0140345P. 11-JUN-1999; Wraight CJ, Werther GA, Edmondson SR;

(MURD-) MURDOCH CHILDRENS RES INST.

VPI; 2001-041421/05.

Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation.

Example 8; Page 62; 201pp; English.

The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an

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           receptor, IgF binding protein [IGPBP] -2 or IGPBP], which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligomucleotide which can be used to design the antisense oligomucleotide which can be used to design the antisense oligomucleotides of the present invention (see AAF45151 and AAF45153 + F45161). The method is useful for ameliorating the effects of psoriasis, inthyosis, pityriasis, ruba, pllaris, serborrhoea, keloids, keratosis, neoplasis, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperpalsia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to an isolated polynucleotide comprising a sequence which is a polymorphic variant of a reference sequence for crystallin, beta B1 (CRYBB1, located on chromosome 22q12.1) gene or their fragment, where the polymorphic variant comprises a CRYBB1 isogene defined by a haplotype from haplotypes 1-16 as given in the specification. Also included are a transgenic non-human animal transformed or transfected
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               with the polymorphic variant, a computer system for storing and analysing polymorphism data for CRYBB1 gene, a genome anthology for the CRYBB1 gene which comprises the defined CRYBB1 isogenes, methods of determining an individuals haplotype or genotype as well as methods of determining the association of a particular haplotype with a disease or trait and a composition comprising at least one genotyping oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human, crystallin beta B1, CRYBB1; chromosome 22q12.1; ophthalmalogical; cataract; allele specific oligonucleotide; ASO; 88; haplotype; genotyping; transgenic animal; PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel polymorphic variants of crystallin, beta B1 useful in studying expression and function of the protein, useful for screening candidate drugs to treat diseases e.g. cataract.
                                                                                                                                                                                                                                                                                                                                                                                    0; Gaps
antisense oligonucleotide, {for Insulin-like Growth Factor [IGF]-1
                                                                                                                                                                                                                                                                                                                                      2.9%; Score 12.4; DB 1; Length 15; 92.9%; Pred. No. 2.9e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kliem SE, Koshy B, Rounds E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR primer #1 for human CRYBB1 gene haplotype PS10.
                                                                                                                                                                                                                                                                                             Sequence 15 BP; 1 A; 3 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 28; Page 31; 94pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GENA-) GENAISSANCE PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAS97386 standard; DNA; 15 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                           112 ACCGCAGCAAGTAC 125
                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 92.9*
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                     14 ACAGCAGCAAGTAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-062253/08.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kazemi A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200185998-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 605
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8,569,669,669,669
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(especially allele-specific oligonucleotides (ASO)) for detecting a polymorphism in the CRYBB1. The isogenes or haplotypes are useful for improving the efficiency and reliability of several steps in the discovery and development of drugs for treating diseases associated with CRYBB1 activity, e.g. cataract. and can also be used by the pharmaceutical research scientist to validate CRYBB1 as candidate target for, and in design of clinical trials of candidate drugs for, treating a specific condition drugs or disease predicted to be associated with CRYBB1 activity. The ASOs are useful as probes and primers, and for assaying a polymorphism in the target region. The present sequence is a pcr primer which amplifies a region of CRYBB1 containing one of 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Detecting a base at a pre-determined position in a nucleic acid molecule, comprises performing primer extension reactions using base-specific detection primers in the presence of a nucleotide-degrading enzyme.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to a method for detecting a base at a predetermined position in a nucleic acid molecule. The method comprises performing primer extension reactions using base specific detection primers, each being specific for a particular base at the predetermined position. The allele-specific (AS) primer extension assay method of the invention is useful for detecting an allele-specific base at a predetermined position in a nucleic acid molecule, for high throughput single nucleotide polymorphism (SNP) analysis, and for detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human, allele-specific base detection; primer extension reaction; base-specific detection primer, allele-specific primer extension assay; AS; high throughput; single mucleotide polymorphism; SNP analysis; mutation detection; genetic variation; allele-specific extension; probe;
                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                       y Match 2.9%; Score 12.4; DB 1; Length 15; Local Similarity 92.9%; Pred. No. 2.9e+02; nes 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                    Sequence 15 BP; 3 A; 4 C; 8 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human genomic DNA p53 SNP AS extension probe #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lundeberg J, Ahmadian A, Nyren P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 2; Page 33; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABX96573 standard; DNA; 15 BP.
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23-FEB-2001; 2001US-00791190.
07-FEB-2002; 2002US-00071926.
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                                                                                                                                                                                                                                                                                                                                                                                                                132 CTGGCCCGCCTGGC 145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                            polymorphic sites
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                                                                                                                                                                                                                                                                                                                            Query Match
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       8888888888888888
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Sequence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;

exemplification of the invention

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The present invention relates to a novel method of producing a stabilised double D loop at a target sequence within a double-stranded nucleic acid. The method involves contacting the double-stranded nucleic acid with an oligonuclectide having a first and second strand with at least a region of complementarity in between them. The first oligonuclectide strand has a region that is complementary to a first strand of the target and binds to the recombinase while the second strand is not bound. The methods, oligonuclectides, compositions and kits are useful for detecting and purifying known nucleic acid targets and for manipulating defined nucleic acid target sequence is an annealing oligonuclectide for Kan- target. This sequence is used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Producing a stabilized double D loop at a target sequence within a double stranded nucleic acid comprises contacting the nucleic acid with an oligonucleotide having a first and second strands with a region of
mutations and genetic variations. The new method solves the deficiencies of previous methods by providing a method of allele-specific extension that allows accurate discrimination between matched and mismatched configurations, as well as reducing or eliminating false positive results observed in prior art. The use of two allele-specific primers increases the sensitivity by a factor of two because signals of two extensions are obtained. The present sequence represents a probe used in the examples of
                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                          2.9%; Score 12.4; DB 1; Length 15; 92.9%; Pred. No. 2.9e+02; rive 0; Mismatches 1; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Detection; purification; double D-loop formation; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                UDS15G annealing oligonucleotide for Kan- target.
                                                                                                                                                           Sequence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Usher MG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 11; Page 48; 99pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rice MC,
                                                                                                                                                                                                                                                                                                                                                                                             35
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                                                                                                                                                                                                                                                                   238 GAGGCTGCTTCCCG 251
                                                                                                                                                                                                                                                                                                                                                                                             AAD48683 standard; DNA; 15
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*tag= a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                    13; Conservative
                                                                                                                             the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-046824/04.
                                                                                                                                                                                                                 Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                             AAD48683;
                                                                                                                                                                                                  Query Match
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The present invention relates to a novel method of producing a stabilised double D loop at a target sequence within a double-stranded nucleic acid. The method involves contacting the double-stranded nucleic acid with an oligonucleotide having a first and second strand with at least a region of complementarity in between them. The first oligonucleotide strand has a region that is complementary to a first strand of the target and binds to the recombinase while the second strand is not bound. The methods, oligonucleotides, compositions and kits are useful for detecting and purifying known nucleic acid targets and for manipulating defined nucleic acid target sequences. The present sequence is an oligonucleotide which is used for determination of optimal oligonucleotide composition for the invention
                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Producing a stabilized double D loop at a target sequence within a double stranded nucleic acid comprises contacting the nucleic acid with an oligonuclectide having a first and second strands with a region of
                                Gaps
                              ö
Query Match

2.9%; Score 12.4; DB 1; Length 15;
Best Local Similarity 92.9%; Pred. No. 2.9e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                 '*tag= a
'note= "Locked nucleic acid (LNA)"
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/note= "Locked nucleic acid (LNA)"
                                                                                                                                                                                                                                                                                         Detection; purification; double D-loop formation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Usher MG;
                                                                                                                                                                                                                                                        Oligo O used for double D-loop formation.
                                                                                                                                                                                                                                                                                                                                                    location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rice MC,
                                                                                                                                                             1672/c
AAD48672 standard; DNA; 15 BP.
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28-SEP-2001; 2001US-0325828P.
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/note= "DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              complementarity in between.
                                                                155 CGGCTTCGACTGGG 168
                                                                                                                                                                                                                                (first entry)
                                                                                         1s cecraceacies 2
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/*tag= c
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                                                                                                                                                                                                     AAD48672;
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AAD48685 standard; DNA; 15 BP

(first entry)

24-PEB-2003

AAD48685;

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The method involves contacting the double-stranded nucleic acid with an oligonucleotide having a first and second strand with at least a region of complementarity in between them. The first oligonucleotide strand has a region that is complementary to a first strand of the target and binds to the recombinase while the second strand is not bound. The methods, oligonucleotides, compositions and kits are useful for detecting and purifying known nucleic acid targets and for manipulating defined nucleic generate neomycin phosphorransferase mutant (Kan-) gene. This sequence is used in the exemplification of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Producing a stabilized double D loop at a target sequence within a double stranded nucleic acid comprises contacting the nucleic acid with an oligonucleotide having a first and second strands with a region of complementarity in between.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to a novel method of producing a stabilised double D loop at a target semience within a double D loop at a target semience within a double D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                       Detection; double D-loop formation; neomycin phosphotransferase;
purification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
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                                                                                                                                                                                                                                                                                                                                     Oligo KLO2 used to generate neomycin phosphotransferase mutant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2.9%; Score 12.4; DB 1; Length 15; 92.9%; Pred. No. 2.9e+02; ive 0; Mismatches 1; Indels
                                         / Match
Local Similarity 92.9%; Pred. No. 2.9e+02;
les 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
            Sequence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rice MC, Usher MG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 10; Page 47; 99pp; English.
                                                                                                                                                                                                                                   ВÞ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-MAR-2001; 2001US-0279146P.
28-SEP-2001; 2001US-0325828P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27-MAR-2002; 2002WO-US009691
                                                                                                                155 CGGCTTCGACTGGG 168
                                                                                                                                                                                                                                   AAD48681 standard; DNA; 15
                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                        15 CGGCTACGACTGGG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kmiec EB, Gamper HB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (UYDE ) UNIV DELAWARE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-046824/04.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      40200279495-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                      Inidentified.
                                                                                                                                                                                                                                                                                                  24-FEB-2003
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Best Local S:
Matches 13
                                                                                                                                                                                                                                                                   AAD48681;
                                                Query Match
                                                                                                                                                                                                 RESULT 609
AAD48681/c
                                                                                Matches
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The present invention relates to a novel method of producing a stabilised double D loop at a target sequence within a double-stranded nucleic acid. The method involves contacting the double-stranded nucleic acid with an oligonuclectide having a first and second strand with at least a region of complementarity in between them. The first oligonuclectide strand has a region that is complementary to a first strand of the target and binds to the recombinase while the second strand is not bound. The methods, oligonuclectides, compestions and kits are useful for detecting and purifying known nucleic acid targets and for manipulating defined nucleic acid target sequence is a locked nucleic acid (IMA) which is used as an annealing oligonuclectide for Kan- target. This sequence is used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                       Producing a stabilized double D loop at a target sequence within a double stranded nucleic acid comprises contacting the nucleic acid with an oligonucleotide having a first and second strands with a region of complementarity in between.
                                                                                                                    Detection; purification; double D-loop formation; locked nucleic acid;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Match 2.9%; Score 12.4; DB 1; Length 15; Local Similarity 92.9%; Pred. No. 2.9e+02; les 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Detection; purification; double D-loop formation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                UR15G annealing oligonucleotide for Kan- target.
                                                                                            KLO2 annealing oligonucleotide for Kan- target.
                                                                                                                                                                                                                                                                                                                                                                                    Usher MG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 11; Page 49; 99pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rice MC,
                                                                                                                                                                                                                                                                                                   27-MAR-2001; 2001US-0279146P.
28-SEP-2001; 2001US-0325828P.
                                                                                                                                                                                                                                                                      27-MAR-2002; 2002WO-US009691.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           155 CGGCTTCGACTGGG 168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15 ceecraceacrees 2
                                                                                                                                                                                                                                                                                                                                                     (UYDE ) UNIV DELAWARE.
                                                                                                                                                                                                                                                                                                                                                                                    Gamper HB,
                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-046824/04.
                                                                                                                                                                                                        WO200279495-A2.
                                                                                                                                                                         Unidentified.
                                                                                                                                                                                                                                         10-OCT-2002.
                                                                                                                                                                                                                                                                                                                                                                                     Kmiec EB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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Matches
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155 CGGCTTCGACTGGG 168

ceecraceacrees 2

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Usher MG;

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The present invention relates to a novel method of producing a stabilised double D loop at a target sequence within a double-stranded nucleic acid. The method involves contacting the double-stranded nucleic acid with an oligonucleotide having a first and second strand with at least a region of complementarity in between them. The first oligonucleotide strand has a region that is complementary to a first strand of the target and binds to the recombinase while the second strand is not bound. The methods, oligonucleotides, compositions and kits are useful for detecting and purifying known nucleic acid targets and for manipulating defined nucleic acid target sequences. The present sequence is an oligonucleotide which is used for double D-loop formation. This sequence is used in the
                                                                                                                                                                                                                                      Producing a stabilized double D loop at a target sequence within a double stranded nucleic acid comprises contacting the nucleic acid with an oligonucleotide having a first and second strands with a region of complementarity in between.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; hammerhead ribozyme; DNAzyme; inozyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; anti-HCV; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
2.9%; Score 12.4; DB 1; Length 15;
Best Local Similarity 92.9%; Pred. No. 2.9e+02;
Matches 13; Conservative 0; Mismatches 1; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Anti-HCV enzymatic nucleic acid substrate sequence #5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                           Example 2; Page 35; 99pp; English.
                                                                                                                                            Rice MC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACD66419 standard; RNA; 15 BP.
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08-JUN-2001; 2001US-00877478.
08-JUN-2001; 2001US-0296876P.
24-OCT-2001; 2001US-0337659P.
05-DEC-2001; 2001US-0337055P.
                       27-MAR-2001; 2001US-0279146P.
28-SEP-2001; 2001US-0325828P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-MAR-2002; 2002WO-US009187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    155 CGGCTTCGACTGGG 168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15 CGCTACGACTGGG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-SEP-2003 (first entry)
                                                                                            (UYDE ) UNIV DELAWARE.
                                                                                                                                               Gamper HB,
                                                                                                                                                                                             WPI; 2003-046824/04.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200281494-A1
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                                                                                                                                               Kmiec EB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           XXCCCCCCCCXXXTTTTTXXXXCCCCCCXXXXXX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to a novel method of producing a stabilised double D loop at a target sequence within a double-stranded nucleic acid. The method involves contacting the double-stranded nucleic acid with an oligonuclectide having a first and second strand with at least a region of complementarity in between them. The first oligonuclectide strand has a region that is complementary to a first strand of the target and binds to the recombinase while the second strand is not bound. The methods, oligonuclectides, compositions and kits are useful for detecting and purifying known nucleic acid targets and for manipulating defined nucleic acid target sequences. The present sequence is an annealing oligonuclectide for Kan- target. This sequence is used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Producing a stabilized double D loop at a target sequence within a double stranded nucleic acid comprises contacting the nucleic acid with an oligonucleotide having a first and second strands with a region of complementarity in between.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Detection; purification; double D-loop formation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 15 BP; 2 A; 6 C; 5 G; 0 T; 2 U; 0 Other;
                                           Location/Qualifiers
1. .15
/*tag= "THER"
/note= "2'-0-methyl nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Usher MG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Oligo N used for double D-loop formation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 11; Page 48; 99pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rice MC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 612
AAD48648/c
ID AAD48648 standard; DNA; 15 BP.
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                                                                                                                                                                                                                                                                                         27-MAR-2002; 2002WO-US009691.
                                                                                                                                                                                                                                                                                                                                    27-MAR-2001; 2001US-0279146P.
28-SEP-2001; 2001US-0325828P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       155 CGGCTTCGACTGGG 168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-FEB-2003 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gamper HB,
                                                                                                                                                                                                                                                                                                                                                                                                            (UYDE ) UNIV DELAWARE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-046824/04.
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                                                                                                                                                                                               WO200279495-A2
                                                Key
modified_base
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Unidentified
                                                                                                                                                                                                                                           10-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kmiec EB,
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Gaps ö 08-JUN-2001; 2001US-00877478. 08-JUN-2001; 2001US-0258876P. 24-OCT-2001; 2001US-0335059P. 05-DEC-2001; 2001US-0337055P.

RIBOZYME PHARM INC

RIBO-)

BLATT L.
MACEJAK D.
MCSWIGGEN J.
MORRISSEY D.
PAVCO P.
LEE P.
DRAPER K. (BLAT/) BLATT L.
(MACE/) MACEJAK D.
(MCSW) MCSHIGGEN J.
(MORK/) MORRISSEY D.
(PAVC/) PAVCO P.
(LEEP/) LEE P.
(LREP/) DRAPER K.
(ROBE/) ROBERTS E.

26-MAR-2002; 2002WO-US009187

WO200281494-A1 17-0CT-2002

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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HEV) or Hepatitis B virus (HEV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, inczymes, zinzymes, amberzymes, and G-cleaver ribozymes. Also disclosed zeroscriptase and/or HEV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV genes and incleic acids may be used to modulate the expression of HBV genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and methods of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene carcinoma. The present sequence represents a substrate for one of the carcinoma. The present sequence espeseents a substrate for one of the carcinoma. The present sequence represents a substrate for one of the
                                                                                                                                                                                        Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
                                                                                                                               Mcswiggen J, Morrissey D, Pavco P, Lee P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 15 BP; 2 A; 6 C; 2 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                          Claim 1; Page 326; 387pp; English.
            RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 92.5.
Best Local 31 Conservative
                                                                                                                                            Roberts E;
                        BLATT L.
MACEJAK D.
MCSWIGGEN J.
MORRISSEY D.
PAVCO P.
                                                                                                                                Macejak D,
                                                                                                                                                                 WPI; 2003-229207/22.
                                                                                                         ROBERTS E.
                                                                                 LEE P.
DRAPER K.
                                                                                                                   Blatt L, Mus
                                                                                                                                                                                                                    .nfection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       invention
           (RIBO-)
(BLAT/)
(MACE/)
(MCSW/)
(MORR/)
(PAVC/)
(LEEP/)
(DRAP/)
%XCCCCCCCCCCCCCXXXX111XX8X11XX8ABBBBBBBBBBBBB
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Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus

Mcswiggen J, Morrissey D, Pavco P, Lee P;

Blatt L, Macejak D, Draper K, Roberts E;

WPI; 2003-229207/22

infection.

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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HRV) or Hepatitis C virus (HCV) or an enzymatic nucleic acids such as harmerhead ribozymes, DNAzymes, and enzymatic nucleic acids such as harmerhead ribozymes, DNAzymes, and enzymatic nucleic acid decoy molecules and G-cleaver ribozymes, Also disclosed are nucleic acid decoy molecules and G-cleaver ribozymes, Also disclosed transcriptase and/or HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and chose the invention and dofor replication of HCV. The compounds and disease states related to HBV and HCV infection, replication and generative and disease states related to HBV and HCV infection, replication and generations. The present sequence represents a target for one of the anti-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Microarray; probe; Mycobacterium; antibiotic-resistance; genotyping; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
2.9%; Score 12.4; DB 1; Length 15;
Best Local Similarity 92.9%; Pred. No. 2.9e+02;
Matches 13; Conservative 0; Mismatches 1; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 15 BP; 2 A; 6 C; 2 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mycobacterium gastrii specific probe GAS-03.
Claim 1; Page 322; 387pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ACC73353 standard; DNA; 15 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-JUL-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ACC73353;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACC73353/
ID ACC7
XX
AC ACC7
XX
DT 15-J
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XX
XX
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0; Gaps

2.9%; Score 12.4; DB 1; Length 15; 92.9%; Pred. No. 2.9e+02; artive 0; Mismatches 1; Indels

6 GGAGTGAAACTGCG 19

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Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; harmerihead ribozyme; DNAzyme; inozyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; anti-HCV; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antihifammatory; target; ss.

Hepatitis C virus

Anti-HCV nucleic acid molecule target sequence #232.

23-SEP-2003 (first entry)

ACD66349;

ACD66349 standard; RNA; 15 BP.

RESULT 614 ACD66349/c

17-APR-2003

Kim C,

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The present sequence is that of kan klo3, an oligonucleotide mismatched (non-hybridising) to the triplet repeat region of exon 1 of the human thurtington's disease (HD) gene. The oligonucleotide is modified by including locked nucleic acid (INA) residues at both ends. Administration of this short, modified oligonucleotide to neuronal PC12 cells bearing an protein (huntingtin) aggregation in cell culture studies. The invention relates to oligonucleotides, including oligonucleotides containing LNA modifications, that alter the genomic HD gene sequence and/or reduce the propensity of huntingtin to form intracellular aggregates. These can be used for the treatment or prevention of HD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New single stranded oligonucleotides comprising a DNA domain having at least one mismatch with respect to the genetic sequence of the Huntington's disease gene to be altered, useful for treating or preventing Huntington's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.9%; Score 12.4; DB 1; Length 15; 92.9%; Pred. No. 2.9e+02; tive 0; Mismatches 1; Indels
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'note= "locked nucleic acid"
                                                                         note = "locked nucleic acid'
                                                                                                           *tag= e
mod_base= OTHER
note= "locked nucleic acid"
                                                                                                                                                                                 *tag= f
|mod_base= OTHER
|note= "locked nucleic acid"
                                                                                                                                                                                                                                                                                                  /note= "locked nucleic acid"
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                                                                                                                                                                                                                                                                /*tag= g
/mod_base= OTHER
                                                       'mod_base= OTHER
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08-AUG-2001; 2001US-0310889P.
04-DEC-2001; 2001US-0337219P.
                                                                                                                                                                                                                                                                                                                                                                                                                  07-AUG-2002; 2002WO-US025352.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kmiec EB, Parekh-Olmedo H;
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15 CGGCTACGACTGGG 2
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Best Local Similarity 92.9°
Matches 13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a microarray comprising a support, a first probe for genotyping Mycobacterium species, second probe for differentiating Mycobacterium tuberoulosis strains, and a third probe for detecting antibiotic-resistant strains, where the probes are immobilized on the support. This sequence represents an example of the first probe used for genotyping Mycobacterium species. The array is useful for simultaneously genotyping Mycobacterium species, differentiating M. tuberculosis strains and detecting antibiotic-resistant strains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                              Microarray for simultaneously genotyping Mycobacteria species, differentiating Mycobacterium tuberculosis strains and detecting antibiotic-resistant strains, comprises specific probes on a support.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Huntington's disease; nootropic; anticonvulsant; huntingtin; human;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              / Match 2.9%; Score 12.4; DB 1; Length 15; Local Similarity 92.9%; Pred. No. 2.9e+02; les 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Locked nucleic acid-containing oligonucleotide kan klo3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 15 BP; 1 A; 3 C; 4 G; 7 T; 0 U; 0 Other;
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/mod_base= OTHER
/note= "locked nucleic acid"
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/mod_base= OTHER
/note= "locked nucleic acid"
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/mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABZ81751 standard; DNA; 15 BP
                                                                                                                                         09-OCT-2002; 2002WO-KR001885.
                                                                                                                                                                           09-OCT-2001; 2001KR-00062125.
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                             Mycobacterium gastrii
                                                                                                                                                                                                                                                                                                                              WPI; 2003-403109/38.
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(KIMC/) KIM C.
(PARK/) PARK H.
                                                                 WO2003031654-A1
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Query Match

Best Loca Matches

8

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Gaps ö

Synthetic

ABZ81751;

RESULT 616 ABZ81751

Locked nucleic acid-containing oligonucleotide kan klo2.

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Huntington's disease; nootropic; anticonvulsant; huntingtin; human; locked nucleic acid; gene therapy; ss.
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/mod_base= OTHER
/note= "locked nucleic acid"
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mod_base= OTHER
'note= "locked nucleic acid"
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/mod_base= OTHER
/note= "locked nucleic acid"
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'mod_base= OTHER
'note= "locked nucleic acid"
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/mod_base= OTHER
/note= "locked nucleic acid"
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/mod_base= OTHER
/note= "locked nucleic acid"
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/mod_base= OTHER
/note= "locked nucleic acid"
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|mod_base= OTHER
|note= "locked nucleic acid"
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mod_base= OTHER
fnote= "locked nucleic acid"
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mod_base= OTHER
note= "locked nucleic acid"
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mod_base= OTHER
/note= "locked nucleic acid"
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/mod_base= OTHER
/note= "locked nucleic acid"
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mod_base= OTHER
note= "locked nucleic acid"
                                                              Location/Qualifiers
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/mod_base= OTHER
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'mod_base= OTHER
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                                      Homo sapiens.
Synthetic.
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                                                                                                                                                                                                          New single stranded oligonucleotides comprising a DNA domain having at least one mismatch with respect to the genetic sequence of the Huntington's disease gene to be altered, useful for treating or preventing Huntington's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Huntington's disease gene non-specific oligonucleotide Kan uD7T/15G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Huntington's disease; nootropic; anticonvulsant; phosphorothioate; huntingtin; human; gene therapy; ss.
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2.9%; Score 12.4; DB 1; Length 15;
Best Local Similarity 92.9%; Pred. No. 2.9e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
1..15
1.kag= a
/mcd base= OTHER
/note= "phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                         Example 5; Page 71; 133pp; English.
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                                      07-AUG-2001; 2001US-0310757P.
08-AUG-2001; 2001US-0310770P.
08-AUG-2001; 2001US-0310889P.
04-DEC-2001; 2001US-0337219P.
           07-AUG-2002; 2002WO-US025352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        155 CGGCTTCGACTGGG 168
                                                                                                                                                   Kmiec EB, Parekh-Olmedo H;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15 CGCTACGACTGGG 2
                                                                                                                  (UYDE ) UNIV DELAWARE
                                                                                                                                                                               WPI; 2003-256478/25.
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modified_base
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ABZ81742/c
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The present sequence is that of single-stranded oligonucleotide Kan MuklSG, which has 2'0-Me modifications throughout its length.

Administration of this oligonucleotide to PCI2 neuronal cells containing an engineered Huntington's disease (HD) gene exon 1 including alternating, repeating Gln codons (CAA/G) had little effect on HD protein (huntingtin) aggregation. This was in contrast to other modified oligonucleotides (see ABZB173'-39) which, although non-specific and non-hybridising to the HD gene, and being incapable of directing sequence hybridising of the tripler repeat region of exon 1, nevertheless reduced the formation of HD protein aggregates. Such oligonucleotides can be used for the treatment or prevention of HD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Distinguishing nonsupercoiled target nucleic acid in sample of nucleic acids from variants of the target, by forming deproteinization-stable double D loops in target query region which distinguish target from
                                                                                                                                                 New single stranded oligonucleotides comprising a DNA domain having at least one mismatch with respect to the genetic sequence of the Huntington's disease gene to be altered, useful for treating or preventing Huntington's disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nonsupercoiled nucleic acid; target query region; genotyping; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                   Example 1; Page 59; 133pp; English.
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                                         Parekh-Olmedo H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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                                                                                        WPI; 2003-256478/25.
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                                         Kmiec EB,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence is that of single-stranded phosphorothioate oligonucleotide Kan ub77/15G, Administration of this oligonucleotide to oligonucleotide collegues of the soligonucleotide to PC12 neuronal cells containing an engineered Huntington's disease (HD) sense exon including alternating, repeating Gln codons (CAA/G) resulted in a reduction in the formation of HD protein (Muntingtin). Kan ub77/15G is an example of modified oligonucleotides of the invention, which although non-specific and non-hybridising to the HD gene, and incapable of directing sequence alternation of the triplet repeat region of exon 1, nevertheless reduce the formation of HD protein aggregates. Such oligonucleotides can be used for the treatment or prevention of HD
                                                                                                                                                                                                                                                                                       New single stranded oligonucleotides comprising a DNA domain having at least one mismatch with respect to the genetic sequence of the Huntington's disease gene to be altered, useful for treating or preventing Huntington's disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Huntington's disease gene non-specific oligonucleotide Kan uR/15G.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 60; 133pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /mod base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABZ81741 standard, RNA; 15 BP.
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08-AUG-2001; 2001US-031070P.

08-AUG-2001; 2001US-031089P.

04-DEC-2001; 2001US-0337219P.
08-AUG-2001; 2001US-0310770P.
08-AUG-2001; 2001US-0310889P.
04-DEC-2001; 2001US-0337219P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 92.9
Matches 13; Conservative
                                                                                                                                                                        Kmiec EB, Parekh-Olmedo
                                                                                                                  (UYDE ) UNIV DELAWARE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (UYDE ) UNIV DELAWARE
                                                                                                                                                                                                                                    WPI; 2003-256478/25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gene therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO2003013437-A2
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modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
Synthetic.
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ABZ8
AAC ABZ8
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AXX ABZ8
AXX Hunt
AXX Hunt
AXX Hunt
AXX Homo
OS Homo
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AXX SPT
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AXX WO20
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target by a nucleotide within a common target query region (TQR), traylving using a recombinase to mediate formation of deproteinizationstable by a nucleotide within a common target query region (TQR), stable double D loop in TQR and then distinguishing degree of formation of double D loops that are stable to deproteinization. The method is useful for distinguishing the presence of a nonsupercoiled target mucleic acid such as a linear duplex DNA, covalently closed circle, or artificial chromosome from the presence of nonsupercoiled target variants within a sample of nucleic acids and is also useful for sample of nucleic acids and is also useful for separating a nonsupercoiled double- stranded nucleic acids within a semple of nucleic acids within a sample of nucleic acids within a separating a nonsupercoiled double- stranded nucleic acids within a semple of nucleic acids within a sample of nucleic acids within a simple may be adapted to a variety of existing detection systems, and permit target amplification without PCR, increasing fidelity. The ability to separate desired double stranded targets with allelic simple separate desired double stranded targets with allelic sequence is an oligonucleotide of the invention. Offers
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Seguence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;

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0; Gaps
/ Match
2.9%; Score 12.4; DB 1; Length 15;
Local Similarity 92.9%; Fred. No. 2.9e+02;
les 13; Conservative 0; Mismatches 1; Indels
                                                 1; Indels
                                                    Matches
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155 CGGCTTCGACTGGG 168 15 cecraceacrees 2 à

ADC13793 standard; DNA; 15 BP (first entry) 18-DEC-2003 ADC13793; RESULT 621 ADC13793/c 

Oligonuclectide of the invention #38.

nonsupercoiled nucleic acid; target query region; genotyping; ss.

Synthetic.

WO2003027640-A2.

33-APR-2003.

27-SEP-2002; 2002WO-US031073.

28-SEP-2001; 2001US-0325828F. 27-MAR-2002; 2002WO-US009691.

(UYDE ) UNIV DELAWARE.

WPI; 2003-371937/35 Kmiec EB,

Distinguishing nonsupercoiled target nucleic acid in sample of nucleic acids from variants of the target, by forming deproteinization-stable double D loops in target query region which distinguish target from

Example 10; SEQ ID NO 38; 179pp; English.

The present invention relates to distinguishing presence of a nonsupercoiled target nucleic acid from presence of nonsupercoiled target variants within a sample of nucleic acids, the variants differing from target by a nucleotide within a common target query region (TQR), involving using a recombinase to mediate formation of deproteinization-

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ctable double D loop in TQR and then distinguishing degree of formation of double D loops that are stable to deproteinization. The method is useful for distinguishing the presence of a monsupercoiled target nucleic acid such as a linear duplex DNA, covalently closed circle, or artificial chromosome from the presence of nonsupercoiled target variants within a sample of nucleic acids. The method distinguishes several mosupercoiled targets within the sample of nucleic acids and is also useful for separating a nonsupercoiled double- stranded nucleic acids within a sample of nucleic acids within a sample of nucleic acids within a sample of nucleic acids within a larget of nucleic acids within a sample of nucleic acids within a single sample, may be adapted to a variety of existing detection systems, and permit target amplification without PCR, increasing fidelity. The ability to separate desired double stranded targets with allelic selectivity, with or without contemporaneous detection, offers sequence is an oligonucleotide of the invention.
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Sequence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;

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Gaps
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1 2.9%; Score 12.4; DB 1; Length 15; Similarity 92.9%; Pred. No. 2.9e+02; 13; Conservative 0; Mismatches 1; Indels
                                   13, Conservative
 Query Match
Best Local S
                                     Matches
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ADC13760/c ID ADC13760 standard; DNA; 15 ADC13760; RESULT 622 

Oligonucleotide of the invention #5. 18-DEC-2003 (first entry)

nonsupercoiled nucleic acid; target query region; genotyping; ss.

Synthetic.

WO2003027640-A2.

03-APR-2003.

27-SEP-2002; 2002WO-US031073

28-SBP-2001; 2001US-0325828P. 27-MAR-2002; 2002WO-US009691.

(UYDE ) UNIV DELAWARE.

Kmiec EB, Rice MC;

WPI; 2003-371937/35.

Distinguishing nonsupercoiled target nucleic acid in sample of nucleic acids from variants of the target, by forming deproteinization-stable double D loops in target query region which distinguish target from variant.

Example 2; SEQ ID NO 5; 179pp; English.

The present invention relates to distinguishing presence of a nonsupercoiled target nucleic acid from presence of nonsupercoiled target variants within a sample of nucleic acid, the variants differing from target by a nucleotide within a common target query region (TQR), involving using a recombinase to mediate formation of deprotainization-stable double D loop in TQR and then distinguishing degree of formation of double D loops that are stable to deprotainization. The method is useful for distinguishing the presence of a nonsupercoiled target nucleic

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acid such as a linear duplex DNA, covalently closed circle, or artificial chromosome from the presence of nonsupercoiled target variants within a sample of nucleic acids. The method distinguishes several nonsupercoiled targets within the sample of nucleic acids and is also useful for separating a nonsupercoiled double stranded nucleic acid target from other nonsupercoiled double stranded nucleic acid target from where 10-10000 fold purification is effected. The methods are readily where 10-10000 fold purification is effected. The methods are readily single sample, may be adapted to a variety of existing detection systems, and permit target amplification without PCR, increasing fidelity. The selectivity, with or without contemporaneous detection, offers significant advantages over current genotyping methods. The present sequence is an oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
         889999999999988
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Query Match 2.9%; Score 12.4; DB 1; Length 15; Best Local Similarity 92.9%; Pred. No. 2.9e+02; Matches 13; Conservative 0; Mismatches 1; Indels 155 CGGCTTCGACTGGG 168

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Gaps

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15 CGGCTACGACTGGG 2 ઠે g

ADC13784 standard; DNA; 15 BP (first entry) 18-DEC-2003 ADC13784; RESULT 623 ADC13784/ 

Oligonucleotide of the invention #29.

nonsupercoiled nucleic acid; target query region; genotyping; ss.

Synthetic.

WO2003027640-A2.

3-APR-2003

27-SEP-2002; 2002WO-US031073.

28-SEP-2001; 2001US-0325828P. 27-MAR-2002; 2002WO-US009691.

UYDE ) UNIV DELAWARE

Kmiec EB, Rice MC;

WPI; 2003-371937/35

Distinguishing nonsupercoiled target nucleic acid in sample of nucleic acids from variants of the target, by forming deproteinization-stable double D loops in target query region which distinguish target from variant.

Example 5; SEQ ID NO 29; 179pp; English

The present invention relates to distinguishing presence of a nonsupercoiled target nucleic acid from presence of nonsupercoiled target variants within a sample of nucleic acids, the variants differing from target by a nucleotide within a common target query region (TQR), involving using a recombinase to mediate formation of deproteinization-stable double D loop in TQR and then distinguishing degree of formation of double D loops that are stable to deproteinization. The method is useful for distinguishing the presence of a nonsupercoiled target nucleic card such as a linear duplax DNA, covalently closed circle, or artificial chromosome from the presence of nonsupercoiled target variants within a sample of nucleic acids. The method distinguishes several nonsupercoiled

The present invention relates to distinguishing presence of a nonsupercoiled target nucleic acid from presence of nonsupercoiled target variants within a sample of nucleic acids, the variants differing from target by a nucleotide within a common target query region (TQR), involving using a recombinase to mediate formation of deprotainizationstable double D loop in TQR and then distinguishing degree of formation of double D loops that are stable to deproteinization. The method is useful for distinguishing the presence of a nonsupercoiled target nucleic caid such as a linear duplax DMA, covalently closed circle, or artificial chromosome from the presence of nonsupercoiled target variants within a sample of nucleic acids. The method distinguishes several nonsupercoiled targets within the sample of nucleic acids and is also useful for separating a nonsupercoiled double- stranded nucleic acid target from other nonsupercoiled double- stranded nucleic acid target from other nonsupercoiled double- stranded nucleic acids target from

Distinguishing nonsupercoiled target nucleic acid in sample of nucleic acids from variants of the target, by forming deproteinization-stable double D loops in target from

Example 11; SEQ ID NO 40; 179pp; English.

variant.

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targets within the sample of nucleic acids and is also useful for separating a nonsupercoiled double- stranded mucleic acid target from cher nonsupercoiled double- stranded mucleic acids within a sample of nucleic acids, where 10-10000 fold purification is effected. The methods are readily multiplexed, permitting a large number of loci to be screened within a sample sample, may be adapted to a variety of existing detection systems, and permit target amplification without PCR, increasing fidelity. The selectivity, with or without contemporaneous detection, offers significant advantages over current genotyping methods. The present sequence is an oligonucleotide of the invention.
                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                       nonsupercoiled nucleic acid; target query region; genotyping; ss.
                                                                                                                                                                                     Query Match 2.9%; Score 12.4; DB 1; Length 15; Best Local Similarity 92.9%; Pred. No. 2.9e+02; Matches 13; Conservative 0; Mismatches 1; Indels ; 0;
                                                                                                                                                             Seguence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                             Oligonucleotide of the invention #40.
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                                                                                                                                                                                                                                          155 CGGCTTCGACTGGG 168
                                                                                                                                                                                                                                                                                                                     ADC13795/c
ID ADC13795 standard; DNA; 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
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Gaps

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The present invention relates to distinguishing presence of a nonsupercoiled target nucleic acid from presence of nonsupercoiled target nucleic acids the variants differing from target by a nucleotide within a sample of nucleic acids, the variants differing from target by a nucleotide within a common target query region (TQR). Involving using a recombinase to mediate formation of deproteinization stable double D loops in TQR and then distinguishing degree of formation of double D loops that are stable to deproteinization. The method is useful for distinguishing the presence of a nonsupercoiled target nucleic acid such as a linear duplex DNA, covalently closed circle, or artificial chromosome from the presence of nonsupercoiled target variants within a sample of nucleic acids and is also useful for targets within the sample of nucleic acids and is also useful for separating a nonsupercoiled double- stranded nucleic acid target from the proposed of the sample of nucleic acids within a multiplexed permitting a large number of loci to be greened within a multiplexed permitting a large number of loci to be greened within a
where 10-10000 fold purification is effected. The methods are readily multiplexed, permitting a large number of loci to be screened within a single sample, may be adapted to a variety of existing detection systems, and permit target amplification without PCR, increasing fidelity. The ability to separate desired double stranded targets with allelic selectivity, withour contemporaneous detection, offers significant advantages over current genotyping methods. The present sequence is an oligonucleotide of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nonsupercoiled nucleic acid; target query region; genotyping; ss.
                                                                                                                                                                                                                                                        2.9%; Score 12.4; DB 1; Length 15; 92.9%; Pred. No. 2.98+02; ive 0; Mismatches 1; Indel8
                                                                                                                                                                                                              Sequence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
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27-MAR-2002; 2002WO-US009691.
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                                                                                                                                                                                                                                                                                                                                                   CGGCTTCGACTGGG 168
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                                                                                                                                                                                                                                                                                                         13; Conservative
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                                                                                                                                                                                                                                                                                   Local Similarity
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                                                                                                                                                                                                                                                                                                         Matches
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The invention relates to a novel method for amplifying a DNA using polymerase chain reaction (PCR) comprising synthesising the first region of a base sequence to be amplified by designing a pair of primers so as to place the first region between them and to anneal each other at the 3-end and carrying out a polymerase chain reaction (PCR) using the primers. Subsequently, the second region is synthesised by designing a pair of primers so as to place the second region partly overlapping with the first region of the base sequence between them and to anneal each other at the 3'-end and carrying out a PCR using the primers. Pinally, the first region is annealed to the second region generating the template to carry out a PCR and thus to synthesize a base sequence containing the first and the second regions. The method of the invention may be useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Amplification of a DNA, a gene encoding the repeated sequence of an amino
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                                                                                                                                                                                                                                                                                                                                                             Gaps
and permit target amplification without PCR, increasing fidelity. The ability to separate desired double stranded targets with allelic selectivity, with or without contemporaneous detection, offers significant advantages over current genotyping methods. The present sequence is an oligonucleotide of the invention.
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                                                                                                                                                                                                                                                                                      2.9%; Score 12.4; DB 1; Length 15; 92.9%; Pred. No. 2.9e+02; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR; DNA amplification; ds; mucin-box; G cassette.
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                                                                                                                                                                                                                      Seguence 15 BP; 2 A; 6 C; 5 G; 0 T; 2 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          for amplifying a DNA sequence. The current se
box encoding G cassette DNA of the invention.
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Best Local Similarity 92...
Best Local Similarity
The state of the stat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15 CGGCTACGACTGGG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-375838/36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  JP2002315583-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 acid sequence,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-JAN-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADD68648;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 626
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADD68648
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           88888888
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AAQ57378;

RESULT 627 AAQ57378

Synthetic.

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The inventors claim an oligo-peptide-carrier conjugate in which the three moieties are covalently linked to one another. The peptide provides a cleavable linker which is cleaved by enzymes which do not degrade antisense oligos (ODNs). The ODN-targeting ligand linkage must be stable to serum proteases, yet cleaved by the lyosocmal enzymes in the target cell. The method involves conjugation of an ODN bearing an electrophilic crosslinking gp. to a peptide which bears two mucleophilic gps of differing reactivity. The resulting ODN-peptide conjugate is prepd. to that a nucleophilic handle remains on the peptide. This gp. is used to turher attach the lyosomotropic carrier to the peptide portion of the ODN-peptide conjugate. The peptide is therefore aslo used as a beterobifunctional linker. Two different model ODNs were used - ODNI and connective is complementary to the intiation codon region of the RNA transcript for the Hepatitis B surface antigen in Hep3B cells. (Updated on 25-WAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                     New covalently linked conjugates of oligo:nucleotide, peptide and carrier - utilising surfactant, poly:amine or targetting ligand as lyso somotropic drug carrier.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR; primer; amplification; promoter; graminaceous plant; rice; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2.9%; Score 12.4; DB 1; Length 16;
32.9%; Pred. No. 3.4e+02;
ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 16 BP; 3 A; 7 C; 1 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nucleotide seguence of the RTBV PCR primer 1.
surface antigen; Hep3B cells; ss.
                                                                                                           /*tag= a
/label= H2N-(CH2)6-O-PO2-
/note= "modified site"
                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 19; 77pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rice tungro bacilliform virus.
                                                                                                                                                                                                                                                                                                                                                                                  Reed MW;
                                                                                                                                                                                                                                                                  93WO-US012246
                                                                                                                                                                                                                                                                                                     92US-00991199
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    33 TGGGACGAAGATGG 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
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Les 13; Conservative
                                                                                                                                                                                                                                                                                                                                           (MICR-) MICROPROBE CORP
                                                                                                                                                                                                                                                                                                                                                                                    Gall AA,
                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1994-217541/26
                                                                                                                                                                                                                                                                  15-DEC-1993;
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                                                                                         misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-DEC-1998
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Hepatitis B
                                                                                                                                                                                                                             23-JUN-1994.
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                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                    Meyer RB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This is a ACE mRNA target sequence (nucleotide no. 1771) of an enzymatic RNA molecule (ribozyme) which cleaves mRNA associated with the development or maintenance of a cardiovascular condition. The concn. of the ribozyme necessary to effect a therapeutic treatment is lower than that of an antisense oligomocleotide and the specificity of action is higher. (Updated on 25-WAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                  inhibitor; inhibition; ribozyme; treatment; prevention; psoriasis; asthma; inflammatory diseases; cardiovascular condition; hypertension; arthritis; restenosis; anglotensin converting enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Enzymatic RNA molecules which cleave mRNA - used to treat or prevent inflammatory, arthritic, stenotic or cardiovascular diseases or
                                                                                                                                                                                                                                 Specific; cleavage; target RNA; protein; prophylaxis; expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 12.4; DB 1; Length 16;
Pred. No. 3.4e+02;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5'-hexylamine modified antisense oligo (ODN1).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 16 BP; 5 A; 5 C; 4 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                             Enzymatic RNA molecule ACE mRNA target sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Antisense oligonucleotide; ODN; modified oligo;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 3; Page 23; 65pp; English.
                                                             AAQ57378 standard; mRNA; 16 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             92US-00916763.
92US-00987132.
92US-00989848.
92US-00989849.
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Best Local Similarity 92.9%;
Matches 13; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                               93WO-US006316
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAQ68223 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 craccacrccaace 14
                                                                                                                                      (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Draper KG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1994-048853/06.
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02-MAR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-DEC-1992;
07-DEC-1992;
07-DEC-1992;
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                                                                                                                                                                                                                                                                                                                                                                    WO9402595-A1
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                                                                                                                                      25-MAR-2003
26-JUL-1994
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Gaps

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AAQ68223;

0X2X55X8X8

RESULT 628

ò g AAQ68223,

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This is the nucleotide sequence of a PCR primer used in the amplification of the Rice tungro bacilliform virus (RTBV) promoter. The isolated genome Length transcript promoter from RTBV is used for driving gene expression in the vascular bundles of gramminaceous plants, especially rice, especially where the gene encodes a protein conferring a desired
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New polynucleotides encoding full-length polypeptides, e.g. secretory and/or membrane proteins, useful for developing medicines for diseases in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cytostatic; Anti-inflammatory; Osteopathic; Neuroprotective; Nootropic; Gene Therapy; human; secretory protein; membrane proteins; cancer; inflammatory disease; PCR; primer;
                                                                                                                                                            Rice tungro bacilliform virus promoter - for driving gene expression in vascular bundles of plants.
                                                                                                                                                                                                                                                                                                            Sequence 16 BP; 7 A; 3 C; 2 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human protein-related PCR primer, SEQ ID 3325.
                                                                                                                                                                                                 Disclosure; Col 3; 12pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HELIX RES INST.
RES ASSOC BIOTECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADASS757 standard; DNA; 16.BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-MAR-2002; 2002EP-00006586.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-SEP-2001; 2001JP-00328381.
24-JAN-2002; 2002US-0350435P.
                                     91US-00789738.
                                                            91US-00789738
                                                                                                             Bhattacharyya M;
                                                                                                                                                                                                                                                                                                                                                                                      397 AGAAGGICITCIAC 410
                                                                                                                                                                                                                                                                                                                                                                                                             AGAAGATCTTCTAC 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                              13; Conservative
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                                                                                    UNIW ) UNIV WASHINGTON
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                                                                                                                                    WPI; 1998-582649/49.
                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                    agronimic trait
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                                                            08-NOV-1991;
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Yamamoto J,
             20-OCT-1998,
                                                                                                             RN,
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(REAS-)
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                                                                                                             Beachy
                                                                                                                                                                                                                                                                                                                                                              Matches
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The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VBGF). A partent (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (fL-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, coular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX7575 represent specific examples of nucleic acid molecules from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; trumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor; foetal liver kinase 1; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleic acid molecule modulating VEGF receptor(s) gene expression or stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.
                                                                                                                                                                                                                                                                                Gaps
which the gene is involved, or as target molecules for gene therapy
                                                                     The present invention relates to novel human secretory or membrane proteins (ADA54072-ADA55710) and their coding sequences (ADA52433-ADA54071). The coding sequences are useful in the gene therapy of diseases caused by abnormalies of the proteins, e.g. cancer, inflammatory diseases, osteoporosis or neurological disease. The proteince was used in an example from the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mouse flt-1 VEGF receptor hammerhead ribozyme substrate #647
                                                                                                                                                                                                                                        Query Match
2.9%; Score 12.4; DB 1; Length 16;
Best Local Similarity 92.9%; Pred. No. 3.4e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                        Sequence 16 BP; 2 A; 5 C; 4 G; 5 T; 0 U; 0 Other;
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                                     Example 8; Page 111; 205pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 4; Page 174; 218pp; English.
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AAX75119 Btandard; RNA; 17 BP.
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96US-00584040.
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(CHIR ) CHIRON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                           14 ACAGAACTCGGTGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1997-259017/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-OCT-1995;
11-JAN-1996;
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Gaps

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2.9%; Score 12.4; DB 1; Length 16; 92.9%; Pred. No. 3.4e+02; ative 0; Mismatches 1; Indels

Tamechika I;

Otsuki T, Wakamatsu A, Sato H, Ish Hio Y, Otsuka K, Nagai K, Irie R, Otsuka M, Nagahari K, Masuho Y;

Ishii

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Sequence 17 BP; 1 A; 8 C; 2 G; 0 T; 6 U; 0 Other;
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Gaps ö Score 12.4; DB 1; Length 17; Pred. No. 3.8e+02; 0; Mismatches 1; Indels 2.9%; Query Match
Best Local Similarity 92.2.
Best Local 3; Conservative

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1 GGCCAGGAGTGAAA 14

15 GGCCAGGAGTGAGA 2

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AAZ24186 standard; DNA; 17 BP. RESULT 632 AAZ24186/c 

AAZ24186;

(first entry) 03-FEB-2000 Human BRCA2 primer scorpion B2731 fragment 1.

Detection; genomic DNA variation; inherited disease; microbial infection; hybridisation; primer; 88.

Synthetic.

Homo sapiens

3B2338301-A. 15-DEC-1999 98GB-00025698. 5-NOV-1998;

98GB-00012768. 13-JUN-1998;

Theaker J, Whitcombe DM; (ZENE ) ZENECA LID.

Gibson NJ, Little S,

4PI; 2000-016019/02

Detecting nucleic acids for the diagnosis of heritable genetic disorders and for the detection of microbial organisms in food and biological

Example 7; Page 25; 74pp; English.

This invention describes a novel method (I) for detecting nucleic acids using novel primers and an integrated signaling system. (I) may be used for the detection of variations genomic DNA samples (e.g. from humans, animals and plants). It is particularly useful for detecting inherited diseases (by detecting abnormalities in DNA from patients) and microbial infections (e.g. human immundeficiency virus (HIV) and Hepatitis C infections (e.g. human immundeficiency virus (HIV) and Hepatitis C viruses or bacterial infections of food). (I) provides high levels of sequence specificity, detection sensitivity and high rates of signal amplicity and allowing enhanced specificity based on the ready availability of a target binding region (TargBR) for hybridization with product is the target species so the output signal obtained is directly related to the amount of extended primer. (I) is not dependent on additional hybridization events or enzymatic steps intra- and inter-clarand competition for the probe site is limited so the probe design is simplified and probes which fail to bind under standard assay conditions in separate probe formats may function in (I). Additionally, homogenous assay formats may be derived from (I). Finally, as the interaction is unimplement, the signal reaction is very rapid, permitting increased to the interaction is very rapid, permitting increased in the method of

Sequence 17 BP; 5 A; 1 C; 8 G; 3 T; 0 U; 0 Other;

DB 1; Length 17; 2.9%; Score 12.4;

Query Match

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                                                                                                                              Detection, genomic DNA variation, inherited disease, microbial infection;
hybridisation, primer, ss.
                                                                                                                                                                                                                                                                            Detecting nucleic acids for the diagnosis of heritable genetic disorders and for the detection of microbial organisms in food and biological samples.
       Gaps
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       Indels
Pred. No. 3.8e+02;
0; Mismatches 1;
                                                                                                                                                                                                                                                  Theaker J, Whitcombe DM;
                                                                                                                  Human BRCA2 quencher primer B4249.
                                                                        ВЪ
                                                                                                                                                                                                        98GB-00025698.
                                                                                                                                                                                                                      98GB-00012768
92.98;
                                                                        AAZ24188 standard; DNA; 17
                      74
                                                                                                    (first entry)
Best Local Similarity 92.9
Matches 13; Conservative
                                 16 ACTCTCTGCACTAC
                      61 AGTCTCTGCACTAC
                                                                                                                                                                                                                                                   Gibson NJ, Little S,
                                                                                                                                                                                                                                                                 WPI; 2000-016019/02.
                                                                                                                                                                                                                                     (ZENE ) ZENECA LID.
                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                        25-NOV-1998;
                                                                                                                                                                                                                       13-JUN-1998;
                                                                                                     03-FEB-2000
                                                                                                                                                                            GB2338301-A.
                                                                                                                                                                                           15-DEC-1999
                                                                                                                                                      Synthetic
                                                                                      AAZ24188;
                                                          RESULT 633
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Example 7; Page 26; 74pp; English.

This invention describes a novel method (I) for detecting nucleic acids using novel primers and an integrated signaling system. (I) may be used using novel primers and an integrated signaling system. (I) may be used for the detection of variations genomic DNA samples (e.g. from humans, and plants). It is particularly useful for detecting inherited diseases (by detecting abnormalities in DNA from patients) and microbial confections (e.g. human immunodeficiency virus (HIV) and Hepatitis C infections (e.g. human immunodeficiency virus (HIV) and Hepatitis C viruses or bacterial infections of food). (I) provides high levels of sequence specificity, detection sensitivity and high rates of signal apparance. Only a single detector/primer species is required (improving simplicity and allowing enhanced specificity based on the ready availability of a target binding region (TargBR) for hybridization with product is the target binding region (TargBR) for hybridization with product is the target species so the output signal obtained is directly related to the amount of extended primer. (I) is not dependent on product is the probes which fail to bind under standard assay conditions in separate probe formats may function in (I). Additionally, homogeneous sensely formats may be derived from (I). Finally, as the interaction is unimplement, the signal reaction is very rapid, permitting increased the invention the invention

Sequence 17 BP; 3 A; 8 C; 1 G; 5 T; 0 U; 0 Other;

Gaps ö Query Match
2.9%; Score 12.4; DB 1; Length 17;
Best Local Similarity 92.9%; Pred. No. 3.8e+02;
Matches 13; Conservative 0; Mismatches 1; Indels

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The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NG20). The regulates expression of a neurite growth inhibitor gene (NG20). The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a DNAzyme) an Inozyme (an endolytic nucleic acid cleaving an RNA molecule possessing an NCH motif), a G-cleaver (cleaving RNA with a NXM motif) an amberzyme (cleaving RNA with an NGM triplet), a zinzyme (cleaving RNA with a YGY motif). The CD20-targetting nucleic acid is used to cleave RNA with a YGY motif). The CD20-targetting nucleic acid is used to cleave RNA of CD20 in the presence of a divalent cation that is preferably Mg^22+. The cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more therapies. In particular, the CD20 targetting nucleic acid may be used to treat lymphoma (NHL), bulky low-grade or follicular non-Hodgkin's lymphoma (NHL), bulky low-grade or follicular nucleicalled with (human immunodeficiency virus) associated NHL, mantle-cell
                                                                                                                                                                                                                                                                                                            Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammarhead ribozyme; DNAzyme; inozyme; doleaver; amberzyme; zinzyme; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphoma; leukaemia; human immunodeficiency virus; HIV associated NHL; mantle-cell lymphoma; MCL; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy·induced neuropathy; amyotrophic lateral sclerosis; ALS; Parkinson's disease; ataxia; Huntington's disease; creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury.
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                                                                                                                                                                                                                                                                            Human NOGO Hammerhead Ribozyme #290.
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28-FEB-2000; 2000US-0185516P.
06-MAR-2000; 2000US-0187128P.
                                                                                                                                                  ABK00290 standard; RNA; 17 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RIBO-) RIBOZYME PHARM INC.
12-MAR-2002 (first entry)
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MCSWIGGEN J.
CHOWRIRA B M.
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(MCSW/) 1
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CC lymphoma (MCL), immunocytoma (IMC), small B-cell lymphocytic lymphoma, immune thrombocytopaenia, and inflammatory arthropathy. The NOGO-cc targetting nucleic acid is used to cleave RNA of the NOGO gene in the presence of a divalent cation that is preferably MG²/+. Furthermore, the condition associated with the level of condition associated with the level of condition that in a condition associated with the level of NOGO. The treatment may further comprise the use of one or more therapies. In particular, the NOGO-targetting nucleic acid may be used to therapies. In particular, the NOGO-targetting nucleic acid may be used to therapies. In particular, agreem (CNS) injury and cerebrovascular accident concentral nervous system (CNS) injury and cerebrovascular accident concentration and sease, demential, multiple sclerosis (MS), chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (MS), chemotherapy-induced neuropathy, anyotrophic lateral sclerosis (ALS), chemotherapy-induced neuropathy, and/or other neurodegenerative disease constants which respond to the modulation of NOGO expression. The present constants and an hammerhead ribozyme of the invention Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; Broat; nozyme; d-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphoma; leukaemia; human immunodeficiency virus; HIV associated NHL; mantle-cell lymphoma; MCL; immunocytoma; IMC; immunocytoma; norcke; dementia; inflammatory arbtropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; cerebrovascylar accident; CVA; Alzheimer's disease; multiple sclerosis; parkinson's disease; ataxia; Huntington's disease; creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease. Gaps Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense ö 2.9%; Score 12.4; DB 1; Length 17; 92.9%; Pred. No. 3.88+02; tive 0; Mismatches 1; Indels ( Sequence 17 BP; 3 A; 6 C; 1 G; 0 T; 7 U; 0 Other; Chowrira BM; ABK02397 standard; RNA; 17 BP. 11-FEB-2000; 2000US-0181797P. 28-FEB-2000; 2000US-0185516P. 06-MAR-2000; 2000US-0187128P. 09-FEB-2001; 2001WO-US004273. 288 AAGCTGGTGAAGGA 301 (RIBO-) RIBOZYME PHARM INC. 12-MAR-2002 (first entry) Query Match Best Local Similarity 92.9% Worthes 13, Conservative || ||||||||||| 17 AACTGGTGAAGGA 4 Human NOGO Amberzyme #69. Blatt L, Mcswiggen J, (BLAT/) BLATT L. (MCSW/) MCSWIGGEN J. (CHOW/) CHOWRIRA B M. WPI; 2001-607195/69. WO200159103-A2. Homo sapiens. Synthetic. 16-AUG-2001. ABK02397; RESULT 635 ABK02397 8 ¥86666666666668 ¥8 8 셤

constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury.

Claim 88; Page 132; 200pp; English. 

307 GCCCCGGGGACCGC 320 8 원

ABK01168; ABK01168/

Human NOGO Inozyme #438.

Human; ss; antieense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; DNAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; neu-Hodgkin's lymphoma; NHL; lymphoma; leukaemia; human immunodeficiency virus; HIV associated NHL; mantle-cell lymphoma; MCL; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy·induced neuropathy; amyotrophic lateral sclerosis; ALS; Parkinson's disease; ataxia; Huntington's disease; creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.

Homo sapiens

The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NOGO). The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a nucleic acids an Inozyme of a nucleic acids (e.g. a ribozyme or a nucleic acids an INDM motif), a G-leaver (cleaving RNA with a NRM motif) a G-leaving RNA with a NRM motif) an amberzyme (cleaving RNA with a NRM motif) an amberzyme (cleaving RNA with a NRM motif) and a memberzyme (cleaving RNA with a NGW motif). The CD20-targetting nucleic acid is used to cleave RNA of CD20 in the presence of a divalent cation that is preferably Mg²+. Furthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more therapies. In particular, the CD20 targetting nucleic acid may be used to treat lymphoma, leukamia, becall lymphoma, leukamia, becall lymphoma, (NLI), bulky low-grade or follicular NHI, lymphocytic context lymphoma (NLI), immunocytoma (INC), small B-cell lymphocytic lymphocytic lymphoma (NLI), immunocytoma (INC), small B-cell lymphocytic lymphocytic argetting nucleic acid is used to cleave RNA of the NOGO gene in the presence of a divalent cation that is preferably Mg²+. Furthermore, the nucleic acid may be contacted with a cell to reduce NOGO activity of the nocontercapy-induced neuropathy, the NOGO-targetting nucleic acid may be contacted with a cell to reduce NOGO activity of the contact a patient having a condition associated with the level of therapies. In particular, the NOGO-targetting nucleic acid may be contacted with a cell to reduce NOGO activity of the contact a patient having a condition associated with the level of therapies. In particular, the NOGO-targetting nucleic acid may be contacted with a cell to reduce NOGO activity of the contact system (CNS) injury and cerebrovascular a central n

Sequence 17 BP; 1 A; 9 C; 6 G; 0 T; 1 U; 0 Other;

Query Match
2.9%; Score 12.4; DB 1; Length 17;
Best Local Similarity 92.9%; Pred. No. 3.8e+02;
Matches 13; Conservative 0; Mismatches 1; Indels 0; Gaps

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1 GCCCCGGGGACCCC 14

ABK01168 standard; RNA; 17 BP. (first entry) RESULT 636 

WO200159103-A2. Synthetic. 

16-AUG-2001

39-FEB-2001; 2001WO-US004273

11-FEB-2000; 2000US-0181797P. 28-FEB-2000; 2000US-0185516P. 06-MAR-2000; 2000US-0187128P.

(RIBO-) RIBOZYME PHARM INC. (BLAT/) BLATT L. (MCSW/) MCSWIGGEN J. (CHOW/) CHOWRIRA B M.

Chowrira BM; Blatt L, Mcswiggen J,

WPI; 2001-607195/69.

Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury.

Claim 88; Page 84; 200pp; English.

The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NGO2). The conjustes expression of a neurite growth inhibitor gene (NGO2). The conjustes expression of a neurite growth inhibitor gene (NGO3). The nucleic acids (e.g. a ribozyme or a nucleic acids as a ribozyme or a conjusted an inozyme (aleaving RN with a NRY motif) proposessing an NCH motif), a G-cleaver (cleaving RN with a NRY motif) proposessing an NCH motif), a G-cleaver (cleaving RN with a YGY motif). The CD20-targetLing nucleic acid is used to cleave RNA with a YGY motif). The CD20-targetLing nucleic acid is used to cleave RNA coff CD20 in the presence of a divalent cation that is preferably MG² +. Coff CD20. The treatment may further comprise the use of one or more coff color. The treatment may further comprise the use of one or more coff color. The treatment may further comprise the use of one or more coff color. The treatment may further comprise the use of one or more cleavagenis, and inflammatory arthropathy. The NOGO-color largetting nucleic acid may be contacted with a cell to reduce NOGO gene in the treatment may further comprise the use of one or more correct may be contacted with a cell to reduce NOGO gene in the color of a divalent cation that is preferably MG² +. Furthermore, the nucleic acid may be contacted with a cell to reduce NOGO gene in the color of a divalent cation that is preferably MG² +. Furthermore, the NOGO. The treatment may further comprise the use of one or more color divalence a patient having a condition associated with the level of the color of the cell and treat a patient having a condition associated with the level of the color and cells and divalence a divalence of disease, demented and condition associated with the level of parkingon's disease, ataxia, Huntington's disease, centered to the repond to the modulation of NOGO expression. The present of the analysis of the constr sequence is an inozyme of the invention

Sequence 17 BP; 3 A; 6 C; 1 G; 0 T; 7 U; 0 Other;

; Query Match
2.9%; Score 12.4; DB 1; Length 17;
Best Local Similarity 92.9%; Pred. No. 3.8e+02;
Matches 13; Conservative 0; Mismatches 1; Indels

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Gaps

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RESULT 637 ABK02396

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Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOG0; hammerhead ribozyme; DNAzyme; inozyme; G-dleaver; amberzyme; zinzyme; lymphoma; leukaemia; human; immunodeficiency virus; HIV associated MHL; hymphocytic leukaemia; human; immunodeficiency virus; HIV associated MHL; mantile-cell lymphoma; MCL; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; central nervous system injury; chemetia; chemotherapy-induced neuropathy; amyotrophic leteral sclerosis; chemotherapy-induced neuropathy; amyotrophic leteral sclerosis; ALS; parkinson's disease; multiple sclerosis; harkinson's disease; multiple sclerosis; hearkinson's disease; ataxia; Huntington's disease;
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ABK02396 standard; RNA; 17 BP.
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28-FEB-2000; 2000US-0185516P.
06-MAR-2000; 2000US-0187128P.
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(BLAT/) BLATT L.
(MCSW/) MCSWIGGEN J.
(CHOW/) CHOWRIRA B M.
                                                                                                                                                            12-MAR-2002 (first entry)
                                                                                                                                                                                                                                               Human NOGO Amberzyme #68.
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                                                                                  ABK02396;
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Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury. Claim 88; Page 131; 200pp; English.

The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down control of a neurite growth inhibitor gene (NGOD). The concleic acids may be enzymatic nucleic acid (e.g. a ribozyme or a nucleic acids may be enzymatic nucleic acid cleaving a an RNA molecule CD possessing an NCH mocif,, a G-cleaver (cleaving RNA with a NGH woith) and NGH woith, a zinzwme (cleaving RNA with an NGH woith) a zinzwme (cleaving RNA with an NGN triplet), a zinzwme (cleaving RNA with an PGD20-targetting nucleic acid is used to cleave RNA cc C CD20 in the presence of a divalent cation that is preferably NG²+.

C Furthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more contacted with a cell lymphoma, low-grade or follicular non-contacted with a cell lymphoma, lymphocytic cleaksemia, HIV (human immunodeficiency virus) associated NHL, mantle-cell lymphoma (MCL), immunocytoma (IMC), small B-cell lymphocytic lymphoma, confined and patient cation that is preferably MG²+. Furthermore, the concleave RNA of the NGO gene in the concleace and any beconcated with a cell to reduce of one or more cell and treat a patient having a condition associated with the level of the NGO. The treatment may further comprise the use of one or more

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therapies. In particular, the NOGO-targetting nucleic acid may be used treat central nervous system (CNS) injury and cerebrovascular accident (CVA, stroke), Alzheimer's disease, dementia, multiple sclerosis (MS), chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (MS), Parkinson's disease, ataxia, Huntingcon's disease, Creutzfeldt-Jäköb disease, muscular dystrophy, and/or other neurodegenerative disease states which respond to the modulation of NOGO expression. The present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human GDMLP-1 17-mer scanning SEQ ID NO:4 sequence SEQ ID NO:1012.
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                                                                                                                                                                 Query Match 2.9%; Score 12.4; DB 1; Length 17; Best Local Similarity 92.9%; Pred. No. 3.88+02; Matches 13; Conservative 0; Mismatches 1; Indels 0
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                                                                                                                 sequence is an amberzyme molecule of the invention
                                                                                                                                                Sequence 17 BP; 1 A; 9 C; 6 G; 0 T; 1 U; 0 Other;
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21-SEP-2000; 2000US-0234687P.
27-SEP-2000; 2000US-0236359P.
04-0CT-2000; 2000US-0026359P.
30-JAN-2001; 2001WO-US000661.
30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000669.
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                                                                                                                                                                                                                                               307 GCCCCGGGGACCGC 320
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                                                                                                                                                                                                                                                                               2 GCCCCGGGGACCCC 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200192524-A2.
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New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.

WPI; 2002-179446/23.

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nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for protein variants having desired phenotypic improvements, and for case as immunogens to raise antibodies that specifically recognise may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as characteristic supplement in patients having specific deficiency in hGDMLP-1 production, and in vacatines or for replacement therapy. The production, and in vacatines or for replacement therapy. The production and secletal muscle disorders. HGDMLP-1 is localised to chromosome 22.

The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at fig. wipo.int/pub/published_pot_sequence
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0; Gaps
                                     Ouery Match
2.9%; Score 12.4; DB 1; Length 17;
Best Local Similarity 92.9%; Pred. No. 3.8e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
Sequence 17 BP; 8 A; 1 C; 8 G; 0 T; 0 U; 0 Other;
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Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
                                                                                        Human GDMLP-1 17-mer scanning SEQ ID NO:4 sequence SEQ ID NO:1011.
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21-SEP-2000; 2000US-0234687P.
27-SEP-2000; 2000US-0236559F.
04-OCT-2000; 2000US-0236559F.
30-JAN-2001; 2001WO-US000662.
30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
                     ABN01019 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                  30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000670.
                                                                                                                                                                                                                                                                                                                                                                                                                      05-FEB-2001; 2001US-0266860P
                                                                                                                                                                                                                                 25-MAY-2001; 2001WO-US016981
                                                                   (first entry)
                                                                                                                                                                                    WO200192524-A2.
                                                                 29-MAY-2002
                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                          06-DEC-2001
                                            ABN01019;
RESULT 639
         ABN01019
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Shannon ME;

Chen W,

DR, Rank

DK,

Hanzel

Penn SG,

Ji Y,

Gu Y.

(AEOM-) AEOMICA INC.

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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used an grobes to detect, characterise and quantify nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as specifically recognise hGDMLP-1 proteins, as specifically recognise hGDMLP-1 proteins, as specific biomolecule and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as the requence or surface-enhanced laser desorption indisation, and production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production and with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hospithied muscle data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at fire, wipo.int/pub/published_pct_sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; POSHL 1; SH3 domain; POSH-like signalling protein 1; oncoge:
Rho GTPase; signal transduction; gene expression; cancer; vaccine;
gene therapy; transgenic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
2.9%; Score 12.4; DB 1; Length 17;
Best Local Similarity 92.9%; Pred. No. 3.8e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human POSHL1 scanning oligonucleotide SEQ ID NO 1823.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 8 A; 1 C; 8 G; 0 T; 0 U; 0 Other;
                                                                                                               Disclosure; SEQ ID NO 1011; 214pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-JAN-2001; 2001MO-US000663.
30-JAN-2001; 2001MO-US000664.
30-JAN-2001; 2001MO-US000665.
30-JAN-2001; 2001MO-US000666.
30-JAN-2001; 2001MO-US000668.
30-JAN-2001; 2001MO-US000668.
30-JAN-2001; 2001MO-US000669.
30-JAN-2001; 2001MO-US000669.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABV91110 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-JAN-2002; 2002EP-00001165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                204 GTGAAAGCAGAAA 217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 GGGAAAGCAGAGAA 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-DEC-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EP1239051-A2
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10-OCT-2001; 2001US-0328205P

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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSH1 1) polypeptide (I), comprising a sequence of 730 amino acids (SI, ABBB3999), a sequence having 55 sequence of dentity to (S1), (S1), having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSH1 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (I) and mucleic acids (II) for identifying a specific binding partner. (I) and mucleic acids (II) caused by altered expression of human POSH1 including disease and treating caused by altered expression of human POSH1 including disease and (II) is treating cancer, they useful in the development of vaccines and (II) is treating cancer therapy. (II) is useful for constructing microarrapy which are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples of the invention. Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to between by the European Patent Office
                                                                                                                                                           Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; POSHL 1; SH3 domain; POSH-like signalling protein 1; oncogene; Rho GTPase; signal transduction; gene expression; cancer; vaccine; gene therapy; transgenic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 2.9%; Score 12.4; DB 1; Length 17; Best Local Similarity 92.9%; Pred. No. 3.8e+02; Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                            Example 2; SEQ ID NO 1823; 60pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human POSHL1 scanning oligonucleotide SEQ ID NO 1824.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17 BP; 4 A; 8 C; 3 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABV91111 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-JAN-2002; 2002EP-00001165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              57 GAGGAGTCTCTGCA 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-DEC-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15 GAGGGGTCTCTGCA 2
                                                                                                                        WPI; 2002-684061/74.
                                         (AEOM-) AEOMICA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EP1239051-A2.
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                                                                                   Shannon M;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 641
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Gaps ö

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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (1), comprising a sequence of 730 amino protein 1 (POSHL 1) polypeptide (1), comprising a sequence of 730 amino (C. acids (S1, ABB8399), a sequence having 65% sequence identity to (S1), comprising the sequence comprising at least 8 contiguous amino sor a fragment of the sequence comprising at least 8 contiguous amino acide. Human POSHL 1 is a proto-oncogene/oncogene product that functions as an cadaptor protein that interacts with Rho family small GTPsses as well as commister protein that interacts with Rho family small GTPsses as well as commister of the signal transduction pathway. (I) is useful of or identifying a specific binding partner. (I) and mucleic acids (II) conceding (I) are useful for diagnosable, monitoring disease and treating caused by altered expression of human POSHL1 including diagnosing and cuseful in gene therapy. (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating cuseful for measuring and for surveying gene expression and creating creating cance is sequence is that of a scanning oligonucleotide useful in examples contined specification, but is based on sequence information supplied to contine by the European Patent Office
                                                                                                                                                                                                                Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; POSHL 1; SH3 domain; POSH-like signalling protein 1; oncogene; Rho GTPase; signal transduction; gene expression; cancer; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                Example 2; SEQ ID NO 1824; 60pp + Sequence Listing; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17 BP; 4 A; 7 C; 4 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABV91108 standard; DNA; 17 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
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30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000670.
23-MAY-2001; 2001US-00864761.
10-OCT-2001; 2001US-0328205P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           57 GAGGAGTCTCTGCA 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14 GAGGGTCTCTGCA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-DEC-2002 (first entry)
                                                                                                                                                                                WPI; 2002-684061/74.
                                                                                                     (AEOM-) AEOMICA INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11-SEP-2002
                                                                                                                                             Shannon M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Si
Matches 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABV91108;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 642
ABV91108/c
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30-JAN-2001, 2001MO-US006664. 30-JAN-2001, 2001MO-US006665. 30-JAN-2001, 2001MO-US000666. 30-JAN-2001, 2001MO-US000666. 30-JAN-2001, 2001MO-US000669. 30-JAN-2001, 2001MO-US000669. 30-JAN-2001, 2001MO-US006670. 23-MAY-2001, 2001US-00864761.

(AEOM-) AEOMICA INC

Shannon M;

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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino cids (SI, ABBB3999), a sequence having 65% sequence of 6711, (S1), having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (I) and nucleic acids (II) for identifying a specific binding partner. (I) and nucleic acids (II) caused by altered expression of human POSHL1 including disease and treating caused by altered expression of human POSHL1 including microarraps which are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples of the invention. Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to berwent by the European Patent Office
                                                                                                                                                                                                                                                                                                                     Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1.
                                                                                                                                                                                                                                                                                                                                                                                                                  Example 2; SEQ ID NO 1821; 60pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17 BP; 3 A; 9 C; 3 G; 2 T; 0 U; 0 Other;
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US0006770.
31-JAN-2001; 2001US-008064761.
                                                                                                                                                                                                                                                                             WPI; 2002-684061/74.
                                                                                                                                                                                     (AEOM-) AEOMICA INC
                                                                                                                                                                                                                                    Shannon M;
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0; Gaps
Query Match 2.9%; Score 12.4; DB 1; Length 17; Best Local Similarity 92.9%; Pred. No. 3.8e+02; Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                        57 GAGGAGTCTCTGCA 70
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Human POSHL1 scanning oligonucleotide SEQ ID NO 1822.
        ABV91109 standard; DNA; 17 BP
                          (first entry)
                          23-DEC-2002
                 ABV91109;
643
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Human, POSHL 1; SH3 domain, POSH-like signalling protein 1; oncogene; Rho GTPase; signal transduction; gene expression; cancer; vaccine;

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gene therapy; transgenic; ss.
                               sapiens
                               Ношо
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3P1239051-A2 11-SEP-2002

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28-JAN-2002; 2002EP-00001165
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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino acids (S1, ABBB399), a sequence having 65% sequence identity to (S1), (C) having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful of downstream components of the signal transduction pathway. (I) is useful of contentlying a specific binding partner. (I) and nucleic acids (II) contenting (I) are useful for diagnoshing, monitoring disease and treating caused by altered expression of human POSHL1 including diagnoshing and cruseful in gene therapy. (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligomucleotide useful in examples of prient specification, but is based on sequence information supplied to between by the European Patent Office
                                                                                               Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1.
                                                                                                                                                                                                                                                                                                                                 Example 2; SEQ ID NO 1822; 60pp + Sequence Listing; English.
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WPI; 2002-684061/74.
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Gaps
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Query Match
2.9%; Score 12.4; DB 1; Length 17;
Best Local Similarity 92.9%; Pred. No. 3.88+02;
Matches 13; Conservative 0; Mismatches 1; Indels
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Human HLA genotyping oligonucleotide SEQ ID NO 863.
                                                                              ABL31374(Standard; DNA; 17 BP.
XX
XC
ABL31374;
XX
C
ABL31374;
XX
C
ABL31374;
XX
DE
Human HLA genotyping oligonucleot
XX
KW
Human; human leukocyte antigen; kW
immunogenetic; transplantation; g
XX
XX
XX
XX
PM
WO200192572-A1.
XX
PD
O6-DEC-2001.
57 GAGGAGTCTCTGCA 70
                                                                       RESULT 644
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Human; human leukocyte antigen; HLA; genotype; polymorphism; immunogenetic; transplantation; genetic disease; ss.

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The invention relates to a typing kit for judging human leukocyte antigen (HLA) genotype of a sample by hybridising a substrate on which 10-24 base oligonucleotides (ABLS) 12-ABLS1809) originating in the sequences of genes e.g. belonging to HLA class I antigens on human genome and containing gene polymorphisms as alloantigens have been immobilised as printers for amplification of cleaved nucleic acids relating to gene polymorphisms. The method is useful for judging HLA genotypes of individuals by determining immunogenetic differences before transplanting between them, providing genetic information to decide comparibility of organ and tissue for transplantation e.g. of bone marrow, kidney, liver, pancreas, Langerhans islet in pancreas and cornea, susceptibility diagnosis of genetic diseases and identifying individuals
                                                                                                                                                                                      Human leukocyte antigen (HLA) typing, useful for judging HLA genotypes of individuals e.g. by determining immunogenetic differences when transplanting between them.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New nucleic acid sequences associated with tumor suppression, regression, apoptosis or virus resistance are useful to diagnose and treat viral
                                                                                                                    Ichihara T, Matsumura Y, Moriya S, Nishida
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ss; tumour suppressor; antitumour; cytostatic; tumour suppression; tumour regression; apoptosis; virus resistance; diagnosis; cellular degeneration.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.9%; Score 12.4; DB 1; Length 17; 92.9%; Pred. No. 3.8e+02; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17 BP; 3 A; 3 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human tumour suppressor sequence #2172.
                                                                                                                                                                                                                                                           Claim 10; Page 257; 345pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Amson R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (MOLE-) MOLECULAR ENGINES LAB SA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ACC53405 standard; DNA; 17 BP.
                               01-JUN-2000; 2000JP-00164798.
01-JUN-2001; 2001WO-JP004662.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                60 GAGTCTCTGCACTA 73
                                                                 (NISN ) NISSHINBO IND INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16 GAGTCTCTGCACAA 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 92.9°
Matches 13; Conservative
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                                                                                    SYST-) SYSTEM RES INC.
                                                                                                                      Inoko H, Kagiya T,
                                                                                                                                                        WPI; 2002-122074/16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-DEC-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 645
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Gaps

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The invention relates to a novel isolated 17 mer nucleic acid sequence, given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, as sequence, with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that they indicates to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers for detecting, indentifying, quantifying and/or amplifying a nucleic acid, e.g. as one component of a gene chip, in vitro as (anti) sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, polypeptides, vectors concaining the nucleic acids, cells containing the vector or antibodies directed against the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral
                                                                                                        The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; sense; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies
                                                                              This sequence represents an isolated nucleic acid sequence associated with tumour suppression or regression, apoptocisis or virus resistence. invention relates to these sequences or sequences having at least 80% identity to them, and polypeptides encoded by the sequences or polypeptides having 80% identity to the polypeptide sequences. The invention is used to disapnose or treat viral disease or disease characterized by development of tumour cells or cellular degeneration
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tumour suppression related human fukutin oligo SEQ ID No 4836.
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disease, development of tumor cells and cell degeneration.
                                                                                                                                                                                                                                                       Sequence 17 BP; 5 A; 4 C; 5 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Telerman A, Amson R, Tuijnder M;
                                       Claim 1; Page 542; 798pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABT39199 standard; DNA; 17 BP.
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diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia. Analysis of the expression of the 17 mer nucleic acids in patient samples is useful for diagnosis and/or prognosis of these diseases. The polypeptides can also be used to generate antibodies, and both the polypeptide and antibodies are useful as components of protein chips. The nucleic acid sequences of the invention can be used in genetherapy. This polymucleotide sequence represents a tumour suppression related human fukutin oligonucleotide of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; GG-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colocectal cancer; brain cancer; oesophageal cancer; colocectal cancer; parceatic cancer; cervical cancer; pad and neck cancer; bladder cancer; panceatic cancer; cervical cancer; pad and neck cancer; cancer; panchago, glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitear; resistant cancer; REL-A-specific inhibitor; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatoid arthritis; restenosis; Crohn's disease; asthma; diabetes; rheumatoid arthritis; restenosis; Crohn's disease; obesity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammatory bowel disease; infection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel enzymatic nucleic acid molecules which down regulates expression a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.
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94US-00245466.
94US-00291932.
96US-00777916.
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MCSW/) MCSWIGGEN J.
DRAPP, DRAPER K G.
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15-AUG-1994;
23-DEC-1996;
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regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFRS), where (I) is an inozyme, zinzwe, G-Cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a partient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially Mg^2^+. The enzymatic and antiense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, ossophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, genetiabline or radiation therapy. The enzymatic and antisense nucleic coid molecules are also useful for treating inflammatory disease such as chief and artificial settle for treating inflammatory disease such as the coid molecules are also useful for treating inflammatory disease such as the coid molecules are also useful for treating inflammatory disease such coid molecules are also useful for treating inflammatory disease such as coid molecules are also useful for treating inflammatory disease or cepting alray inflammation, inflammatory bowel disease or cepting alray inflammation, inflammatory bowel disease or cepting alray allergic alray inflammation, inflammatory bowel disease or conficion, gene therapy application, inflammatory bowel disease or conficion alray inflammation, inflammatory bowel disease or conficion alray and mycoardial), inflammatory bowel disease or conficion allergic alrays inflammation, inflammatory bowel disease or conficion and an order or septiments and an order or sequence represents the substrate of a novel enzymatic
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85.7%; Pred. No. 3.8e+02;
ive 1; Mismatches 1; Indels
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94US-00245466.
94US-00291932.
96US-00777916.
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nes 12, Conservative
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18-MAY-1994;
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23-DEC-1996;
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(STIN/) STINCHCOMB D T.

US2002177568-A1 Homo sapiens

28-NOV-2002

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Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.
                    Stinchcomb DT, Mcswiggen J, Draper KG;
                                                                             Claim 3; Page 50; 72pp; English.
(MCSW/) MCSWIGGEN J. (DRAP/) DRAPER K G.
                                  WPI; 2003-340953/32.
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The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFRS), where (I) is an inozyme, zinzyme, G-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat carcer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially Mg^2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or milidizug resistant cancer. The method involves use of other drug cervical, esistant cancer. The method involves use of other drug chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, colorhosphamide, doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, canid molecules are also useful for treating inflammatory disease such as acid molecules are also useful for treating inflammatory disease such a cheumatoid arthritis, restenosis, asthma, Crohn's disease, diabetes, cobestiv, autoimmune disease, lupus, multiple sclerosis, transplant/graft rejection, gene therapy applications, ischaemial. reperfusion injury, central nervous system (CNS) and myocardial), glomerulonephritis, signates and such and processes and service the service the such as a processes and serviced the service acid molecules are entered to an entered the service acid molecules are also useful for the service acid mol infection. This sequence represents the substrate of a novel enzymatic nucleic acid molecule

Sequence 17 BP; 3 A; 4 C; 8 G; 0 T; 2 U; 0 Other;

0; Gaps 2.9%; Score 12.4; DB 1; Length 17; 85.7%; Pred. No. 3.8e+02; ive 1; Mismatches 1; Indels 12; Conservative Query Match Local Matches

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ACA06441 standard; RNA; 17 BP 03-JUN-2003 (first entry) ACA06441; RESULT 649 ACA06441/c 

NFKB sub-unit modulating inozyme substrate #260.

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Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; oceophageal cancer; prometh cancer; brain cancer; cervical cancer; pancreatic cancer; percreatic cancer; percreatic cancer; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; docetaxel; cisplatin; methotrexate; cyclophosphamide; doxorubhi; fluorouracil carboplatin; edatrexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatoid arthritis; restenosis; Crohn; & disease; besity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/garaft rejection; reperfusion injury; glomerulonephritis; allergic alrway inflammation; inflammatory bowel disease; infection; ss.

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regulates expression of a sequence encoding a subunit of muclear factor kappa B (NFKB), where (I) is an inozyme, zinzyme, G-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A ectivity in a cell, for treating a patient having a condition associated with the level of REL-A.

(I) is useful for clearing RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially MG2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, corporate, colorectal, brain, oscophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug theretapy including paclitaxel, docetaxel, cisplatin, methotrexate, chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, conditabline or radiation therapy. The enzymatic and antisense nucleic conditabline or radiation therapy. The enzymatic and antisense mucleic conditabline or radiation therapy. The enzymatic and antisense mucleic conditabline or radiation therapy. The enzymatic and antisense mucleic conditabline or radiation therapy. The enzymatic and antisense mucleic conditabline or radiation therapy. The enzymatic and antisense mucleic conditabline disease, lupus, multiple soleroals, transplant graft rejection, gene therapy applications ischemial/trepartualon injury inflammatory of some relection, gene therapy applications ischemial/trepartualon injury inflammatory for environment env
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2.9%; Score 12.4; DB 1; Length 17;
Best Local Similarity 92.9%; Pred. No. 3.8e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
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94US-00291932.
96US-00777916.
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(MCSW/) MCSWIGGEN J.
(DRAP/) DRAPER K G.
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G-cleaver, amberzyme, cancer; REL-A activity, breast cancer; human; lung cancer; prostate cancer; REL-A activity, breast cancer; human; lung cancer; prostate cancer; cancer; brain cancer; prostate cancer; permed cancer; brain cancer; cancer; carvical cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemocherapy; paclitaxel; docetaxel; cisplatin; methodrexate; cyclophosphamide; doxorubin; fluorouracil carboplatin; edatrexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatoid arthritis; restenosis; Crohn's disease; obesity; isofnamia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammatory bowel disease; infection; ss. inozyme; zinzyme; Enzymatic nucleic acid; nuclear factor kappa B; NFKB;

Homo sapiens.

US2002177568-A1.

28-NOV-2002.

23-MAY-2001; 2001US-00864785.

92US-00987132. 94US-00245466. 94US-00291932. 96US-00777916. .8-MAY-1994; 07-DEC-1992;

STIN/) STINCHCOMB D T.

33-DEC-1996;

(MCSW/) MCSWIGGEN J (DRAP/) DRAPER K G.

Stinchcomb DT, Mcswiggen J, Draper KG;

WPI; 2003-340953/32.

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.

Claim 3; Page 55; 72pp; English.

The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFRS), where (I) is an inozyme, zinzwe, G-Cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially Mg^2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oespecially Mg^2+. The enzymatic and strictal, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, gemeitabline or radiation therapy. The enzymatic and antisense nucleic acid molecules are also useful for treating inflammatory disease such as acid molecules are also useful for treating inflammatory disease such as acid molecules. rheumatoid arthritis, restenosis, asthma, Crohn's disease, diabetes, obesity, autoimmune disease, lupus, multiple sclerosis, transplant/graft rejection, gene therapy applications, ischaemia/reperfusion injury (central nervous system (CNS) and myocardial), glomerulonephritis, seppis, allergic airway inflammation, inflammatory bowel disease or infection. This sequence represents the substrate of a novel enzymatic nucleic acid molecule

Sequence 17 BP; 0 A; 9 C; 7 G; 0 T; 1 U; 0 Other;

Gaps ö 2.9%; Score 12.4; DB 1; Length 17; ilarity 92.9%; Pred. No. 3.88+02; Conservative 0; Mismatches 1; Indels Best Local Similarity Matches 13; Conserva Query Match

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14

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RESULT 651 ACA06442,

BP. ACA06442 standard; RNA; 17 

ACA06442;

(first entry) 03-JUN-2003 NFKB sub-unit modulating inozyme substrate #261.

Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; praint cancer; colorectal cancer; pancreatic cancer; cervical cancer; pancreatic cancer; lymphoma; glioma; multidrug resistant cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxal; docetaxel; cisplatin; methotrexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatorid arthritis; restenosis; Crohn's disease; abstry; ischaemia; pene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection; ss.

Homo sapiens.

US2002177568-A1.

28-NOV-2002.

23-MAY-2001; 2001US-00864785.

92US-00987132. 94US-00245466. 94US-00291932. 96US-00777916. 18-MAY-1994; 15-AUG-1994; 23-DEC-1996; 07-DEC-1992;

STIN/) STINCHCOMB D T. (MCSW/) MCSWIGGEN J. (DRAP/) DRAPER K G. DRAPER K G. Draper KG; Stinchcomb DT, Mcswiggen J,

WPI; 2003-340953/32.

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.

Claim 3; Page 31; 72pp; English.

The invention describes an enzymatic nucleic acid molecule (1) which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFFD), where (1) is an inozyme, zinzyme, G-Cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in: a cell, for treating a patient having a condition associated with the level of REL-A. (1) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially MG^2+. The enzymatic and nisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, oervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or therapy including paclitaxal, docetaxel, cisplatin, methotrexate, chemotherapy including paclitaxal, docetaxel, cisplatin, methotrexate, cyclohosphamide, doxorubin, fluorouracil carboplatin, distrakate, cyclohosphamide, doxorubin, fluorouracil carboplatin, glioma nucleic acid molecules are also useful for treating inflammatory disease such as the method artificial, restenosis, asthma, Crohn's disease, diabetes,

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lumg cancer; prostate cancer; colosectal cancer; brain cancer; colosectal cancer; brain cancer; colosectal cancer; brain cancer; cervical cancer; stomach cancer; bladder cancer; pancreatic cancer; cervical cancer; stomach cancer; occupant cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxal; docetaxel; cisplatin; methotraxate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatorid arthritis; restenoâis; Crohn's disease; obesity; ischaemia; genc therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection; ss.
obesity, autoimmune disease, lupus, multiple sclerosis, transplant/graf rejection, gene therapy applications, ischaemia/reperfusion injury (central nervous system (CNS) and myocardial), glomerulonephritis, sepsis, allergic airway inflammation, inflammatory bowel disease or infection. This sequence represents the substrate of a novel enzymatic
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                                                                                                                                                                      Query Match 2.9%; Score 12.4; DB 1; Length 17; Best Local Similarity 92.9%; Pred. No. 3.8e+02; Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                       Sequence 17 BP; 1 A; 11 C; 3 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NFKB sub-unit modulating amberzyme substrate #64.
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94US-00245466.
94US-00291932.
96US-00777916.
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                                                                                                                                                                                                                                                      144 GCGGTGGAGGCCGG 157
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                                                                                               nucleic acid molecule
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(DRAP/) DRAPER K G.
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15-AUG-1994;
23-DEC-1996;
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cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A.

(I) is useful for clearing RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially Mc2+. The enzymatic and cantisense mucleic acid molecules are useful for traating breast, lung, prostate, colorectal, brain, oseophageal, stoomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug charpapies unch as monoclonal antibodies, REL-A specific inhibitors or cherapies use and antibodies, REL-A specific inhibitors or cherapy including paclitaxel, docetaxel, cisplatin, edatrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide doxorubin, fluorouracil carboplatin, edatrexate, candidation therapy. The enzymatic and antisense nucleic acid molecules are also useful for treating inflammatory disease such a chematoid arthritis, restenosis, suchaemia/reperfusion injury celection, gene therapy applications, isofaemia/reperfusion injury celection, gene therapy applications, isofaemia/reperfusion injury celection, gene therapy applications, isofaemia/reperfusion injury cepticion. This sequence represents the substrate of a novel enzymatic conception acid molecule
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Pred. No. 3.8e+02;
1; Mismatches 1; Indels
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94US-00245466.
94US-00291932.
96US-00777916.
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Best Local Similarity 85.7%;
Matches 12; Conservative 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      266 GCACCTGGAGCAGG 279
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MCSWIGGEN J.
DRAPER K G.
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18-MAY-1994;
15-AUG-1994;
23-DEC-1996;
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(DRAP/)
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ACA09050/
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Gu Y, Nguyen C;

Shannon M,

WPI; 2003-423107/40.

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Stinchcomb DT, Mcswiggen J, Draper KG;
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                                                                                                                            (first entry)
     WPI; 2003-340953/32
                                                                                                                                                   Homo sapiens.
                                                                                                                                                       3P1281758-A2.
                                                                                                                            20-NOV-2003
                                                                                                                        ADB00481;
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ADB00481/c
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The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor regulates expression of a sequence encoding a subunit of nuclear factor cancer and is useful for down-regulating REI-A activity in a cell, for cancer and is useful for down-regulating REI-A activity in a cell, for creating a patient having a condition associated with the level of REI-A. (I) is useful for cleaving RNA comprising a sequence of REI-A gene, in the presence of a divalent cation, especially Mg^2+ The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or unlittury resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REI-A-specific inhibitors or chemotherapy including paciltaxel, docetaxel, cisplatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexace, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexace, cyclophosphamide, doxorubin, fluorouracil carboplatin, datrexace, cyclophosphamide, doxorubin, gone therapy applications, ischaemia/reperfusion injury (central nervous system (CMS) and myocardial), glomerulonephritis, cspsis, allergic airway inflammation, inflammatory bowel disease or infection. This sequence represents the substrate of a novel enzymatic inflection. This sequence represents the substrate of a novel enzymatic Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases. Claim 3; Page 55; 72pp; English

0; Gaps 2.9%; Score 12.4; DB 1; Length 17; 92.9%; Pred. No. 3.8e+02; ive 0; Mismatches 1; Indels Seguence 17 BP; 0 A; 8 C; 8 G; 0 T; 1 U; 0 Other; 2.9% Best Local Similarity 92.9° Matches 13; Conservative

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ADB00481 standard; DNA; 17 BP 305 GAGCCCCGGGGACC 318 15 GAGCCCCGGGGCCC 2

Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss. Human MDZ3 scanning oligonucleotide SEQ ID 1467.

30-JUL-2002; 2002EP-00016874.

(AEOM-) AEOMICA INC

New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.

Gu Y, Nguyen C;

Shannon M,

WPI; 2003-423107/40.

The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ12, MDZ12. MDZ3 is conceded at chromosome 7422.1, MDZ4 is encoded at chromosome 6721.3-22.2, MDZ7 is encoded at chromosome 6721.3-22.2, MDZ7 is encoded at chromosome 6721.3 and MDZ12 is encoded at chromosome 5721.2 and MDZ12 is encoded at chromosome secretarial and manufacturing a medicament for treating or preventing a disorder or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The probes are also be used as probes to detect and characterize gross alterations in MDZ3, MDZ7, or MDZ12 genetic locus. The probes are brotain are useful at therapeutic agents for gene therapy or as proteins are useful as therapeutic agents for gene therapy or as proteins are useful. ö Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MD21; MD24; MD212; MD212; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss. New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MD23, MD24, MD27 or MD212, e.g. cancer. ö 2.9%; Score 12.4; DB 1; Length 17; 92.9%; Pred. No. 3.8e+02; tive 0; Mismatches 1; Indels Sequence 17 BP; 3 A; 7 C; 5 G; 2 T; 0 U; 0 Other; Human MDZ3 scanning oligonucleotide SEQ ID 1469. Example 8; SEQ ID NO 1467; 103pp; English. ADB00483 standard; DNA; 17 BP. 30-JUL-2002; 2002EP-00016874. 02-AUG-2001; 2001US-00922181 262 CGGTGCACCTGGAG 275 20-NOV-2003 (first entry) 2.9% Query Match Best Local Similarity 92.99 Matches 13, Conservative 17 CGGTGCACCTGCAG 4 (AEOM-) AEOMICA INC. Homo sapiens. 3P1281758-A2 ADB00483; RESULT 655 ADB00483/c THE TEXT TO THE TEXT OF THE TE 8X8X84444X8X8X8X8X8X8X8X8X8X8X8X8 ઠ 셤

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The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ12, MDZ12. MDZ3 is encoded at Chromosome 7422.1, MDZ4 is encoded at Chromosome 6721.3-22.2, MDZ7 is encoded at chromosome 6721.3 and MDZ12 is encoded at chromosome 15q26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or immunifacturing a medicament for treating or preventing a disorder sesociated with Acreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as proteins. The present sequence was used to illustrate the invention.
Example 8; SEQ ID NO 1469; 103pp; English.
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Sequence 17 BP; 3 A; 6 C; 6 G; 2 T; 0 U; 0 Other;

Gaps ö Query Match 2.9%; Score 12.4; DB 1; Length 17; Best Local Similarity 92.9%; Pred. No. 3.8e+02; Matches 13; Conservative 0; Mismatches 1; Indels 13; Conservative

262 CGGTGCACCTGGAG 275 cedrecacerecae 2 15 ઠ 셤

ADA99414 standard; DNA; 17 20-NOV-2003 ADA99414; RESULT 656

BP.

Human MDZ3 scanning oligonucleotide SEQ ID 403. (first entry) 

Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ12; chromosome 7g22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15g26.1; cancer; developmental disorder; ss.

Homo sapiens

35-FEB-2003.

EP1281758-A2.

30-JUL-2002; 2002EP-00016874.

02-AUG-2001; 2001US-00922181.

(AEOM-) AEOMICA INC.

Shannon M, Gu Y, Nguyen C; WPI; 2003-423107/40. New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.

Example 8; SEQ ID NO 403; 103pp; English.

The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7922.1, MDZ4 is encoded at chromosome 6721.3-22.2, MDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 15p11.2 and MDZ12 is encoded at chromosome 15p112.2 and MDZ12 is encoded at chr

or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for disgnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ12 genetic locus. The probes are proteins are useful as thereautic as microarrays for measuring gene expression. The proteins are useful as thereautic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention. %\$3888888888888

Sequence 17 BP; 2 A; 9 C; 1 G; 5 T; 0 U; 0 Other;

ö Query Match 2.9%; Score 12.4; DB 1; Length 17; Best Local Similarity 92.9%; Pred. No. 3.8e+02; Matches 13; Conservative 0; Mismatches 1; Indels

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ADA99489 standard; DNA; 17 BP. ADA99489

ADA99489;

20-NOV-2003

(first entry)

Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss. Human MDZ3 scanning oligonucleotide SEQ ID 478.

Homo sapiens.

EP1281758-A2.

05-FEB-2003.

30-JUL-2002; 2002EP-00016874.

02-AUG-2001; 2001US-00922181. 

(AEOM-) AEOMICA INC

Shannon M, Gu Y, Nguyen

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WPI; 2003-423107/40.

MDZ3 New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MD24, MD27 or MD212, e.g. cancer.

Example 8; SEQ ID NO 478; 103pp; English.

The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MD23, MD24, MD27, MD212. MD23 is encoded at chromosome 7922.1, MD24 is encoded at chromosome for an encoded at chromosome for the modern is encoded at chromosome 15q26.1. The MD21 is encoded at chromosome 15q26.1. The MD24, MD24, MD21, and MD212 is encoded at chromosome is second in manufacturing a medicament for treating or preventing a disorder or in manufacturing a medicament for treating or preventing adjsorder or in massociated with decreased or increased expression or activity of MD23, MD24, MD27, or MD212, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MD23, MD24, MD27, or MD212. The nucleic acids can also be used as probes to detect and characterize gross alterations in MD23, MD24, MD27, or MD212 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The

BP.

ADB00482 standard; DNA; 17

RESULT 659

ADB00482;

293 GGTGAAGGACCTGA 306

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GGTGGAGGACCTGA 14

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The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ12, MDZ12. MDZ3 is proteins and their coding sequences: MDZ3, MDZ4, MDZ12, MDZ12. MDZ3 is cancoded at chromosome f922.1.4, MDZ4 is encoded at chromosome f922.1.2, MDZ7 is encoded at chromosome f922.1.2, mDZ2 is encoded at chromosome f922.1.2 and MDZ12 is encoded at chromosome for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acide and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ1, or MDZ12. The nucleic acide can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as proteins are useful as therapeutic agents for gene therapy or as
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cytostatic; immunostimulant; gene therapy; vaccine; human;
zinc finger protein; MDZ3; MDZ4; MDZ12; chromosome 7q22.1;
chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer;
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                                                                                                                                         Gaps
proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
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                                                                                              ch 2.9%; Score 12.4; DB 1; Length 17; 1 Similarity 92.9%; Pred. No. 3.8e+02; 13; Conservative 0; Mismatches 1; Indels
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                                                            Seguence 17 BP; 3 A; 4 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                        Human MDZ3 scanning oligonucleotide SEQ ID 482.
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                                                                                                                                                                                                                                                                                                                       ADA99493 standard; DNA; 17 BP.
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                                                                                                                                                                                 292 TGGTGAAGGACCTG 305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      developmental disorder; ss
                                                                                                                                                                                                                     (first entry)
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                                                                                                                         Local Similarity
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                                                                                                        Query Match
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The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7q22.1, MDZ4 is encoded at chromosome 7g21.3 MDZ4 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 15p13.2 and MDZ12 is encoded at chromosome 15g26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, 15g26.1. The MDZ3, MDZ4, MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disease associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ1, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, or MDZ12, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as proteins are useful as therapeutic agents for gene therapy or as
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                                                                                                                                                 Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ12; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.
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                                                                                                              Human MDZ3 scanning oligonucleotide SEQ ID 1468.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 8; SEQ ID NO 1468; 103pp; English.
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Best Local Similarity 92.9
Matches 13; Conservative
                                                                           (first entry)
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ADB00484/c
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Gaps

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Query Match 2.9%; Score 12.4; DB 1; Length 17; Best Local Similarity 92.9%; Pred. No. 3.8e+02; Matches 13; Conservative 0; Mismatches 1; Indels

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ADB00484;

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The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-HER2, K-Ras, H-Ras, and HIV acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, shown in NBZS9889 - ABZ62216, ABZ64944 - ABZ65531, ABZ65520 - ABZ65524, ABZ65530 - ABZ65524 - ABZ65531, ABZ65531 - ABZ65524 - ABZ65531 - ABZ65531 - ABZ65524 - ABZ65531 - ABZ65531 - ABZ65534 - ABZ65531 - ABZ65534 - ABZ6554 - 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras; enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV; anti-rheumatic; cancer; AIDS; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 2.9%; Score 12.4; DB 1; Length 17; Best Local Similarity 78.6%; Pred. No. 3.8e+02; Matches 11; Conservative 2; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 4; Page 144; 185pp; English.
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06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
                                                                                                                                                                                                                                                                                                                                                     29-MAY-2002; 2002WO-US016840.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                 Homo sapiens.
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                                                                                                                                                                                                                                                                     Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3,
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                                                                                                                                                                                                                      duman MDZ3 scanning oligonucleotide SEQ ID 1470.
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              ADB00484 standard; DNA; 17 BP
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                                                                                                                                                 (first entry)
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ABZ65139

RESULT 66
ABZ65139
ID ABZ6
XC ABZ6
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DT 21-M
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Gaps

RNA stability, RNA expression, RNA synthesis, antisense, enzymatic nucleic acid, hammerhead ribozyme, DNAzyme, inozyme; zinzyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative, disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; 88.

Nucleic acid molecule, Hepatitis C virus; HCV; Hepatitis B virus; HBV;

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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, inozymes, zinzymes, amberzymes, and G-cleaver ribozymes. Also disclosed are nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV genes and/or HBV verse transcriptate the expression of HBV genes and MBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and methods of the invention are useful for the treatment of degenerative and enthods of the invention are useful for the treatment of degenerative and issues states related to HBV and HCV infection, replication and gene carcinoma. The present sequence represents a substrate for one of the HCV DNAzyme or minus strand DNAzyme sequences disclosed in the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                              Mcswiggen J, Morrissey D, Pavco P, Lee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 2.9%; Score 12.4; DB 1; Length 17; Best Local Similarity 85.7%; Pred. No. 3.8e+02; Matches 12; Conservative 1; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 1; Page 298; 387pp; English.
                                                                                                                       08-JUN-2001; 2001US-00877478.
08-JUN-2001; 2001US-0296876P.
24-OCT-2001; 2001US-0337055P.
05-DEC-2001; 2001US-0337055P.
                                                                      26-MAR-2002; 2002WO-US009187
                                                                                                                                                                                                               RIBOZYME PHARM INC
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                                                                                                                                                                                                                                               MACEJAK D.
MCSWIGGEN J.
MORRISSEY D.
                                                                                                                                                                                                                                                                                                                                                                                              Macejak D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-229207/22
                                                                                                                                                                                                                                                                                                                                                            ROBERTS E.
                                                                                                                                                                                                                                                                                                        PAVCO P.
LEE P.
                                                                                                                                                                                                                                                                                                                                       DRAPER K.
 WO200281494-A1.
                                                                                                        26-MAR-2001;
                                    17-0CT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         infection.
                                                                                                                                                                                                                                                                                                                                                                                                                 Draper K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nvention
                                                                                                                                                                                                                                                                                                                                         (DRAP/)
(ROBE/)
                                                                                                                                                                                                                                                                                                                                                                                                ij
                                                                                                                                                                                                                                                                                   (MORR/)
(PAVC/)
(LEEP/)
                                                                                                                                                                                                                                (BLAT/)
(MACE/)
(MCSW/)
                                                                                                                                                                                                               RIBO-)
                                                                                                                                                                                                                                                                                                                                                                                                Blatt
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26-MAR-2001; 2001US-00917879. 08-UJN-2001; 2001US-00977478. 08-UJN-2001; 2001US-0295676P. 24-OCT-2001; 2001US-0335059P. 05-DEC-2001; 2001US-0337055P.

RIBOZYME PHARM INC. BLATT L.

BLATT 1. MACEJAK D. MCSWIGGEN J. MORRISSEY D.

(RIBO-) (BLAT/) (MACE/) (MCSW/) (MORR/)

26-MAR-2002; 2002WO-US009187

Hepatitis C virus

WO200281494-A1.

17-0CT-2002

ä Lee

Mcswiggen J, Morrissey D, Pavco P,

Macejak D, Roberts E;

Blatt L, N Draper K,

DRAPER K. ROBERTS E.

PAVCO P.

LEE P.

(PAVC/) 1 (LEEP/) 1 (DRAP/) 1 (ROBE/) 1

WPI; 2003-229207/22

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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HWV) DAN The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, an enzymatic nucleic acid such as hammerhead ribozymes, DNAzymes, an uncleic acid decoy molecules and G-cleaver ribozymes. Also disclosed a ranscriptase and/or HBV reverse transcriptase primer sequences, as well as oligomolectides that specifically bind the Enhancer I region of HBV genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and methods of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene carcinoma. The present sequence represente a substrate for one of the HCV DNAZYME or minus strand DNAzyme sequences disclosed in the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
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2.9%; Score 12.4; DB 1; Length 17;
Best Local Similarity 92.9%; Pred. No. 3.8e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 0 A; 6 C; 8 G; 0 T; 3 U; 0.Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; Page 260; 387pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 셤
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0; Gaps

substrate sequence #1477.

BP

RESULT 663

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Sequence 17 BP; 3 A; 8 C; 6 G; 0 T; 0 U; 0 Other;

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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and erzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, or inozymes, zinzymes, amberzymes, and G-cleaver ribozymes, DNAzymes, are nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV DNA. The nucleic acids may be used to modulate the expression of HBV genes and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and methods of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene carcinoma. The present sequence represents a substrate for one of the HCV bindaryme or minus strand DNAzyme sequence disclosed in the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
                                                                                                                                                                 Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; harmerthead ribozyme; DNAzyme; inozyme; zinzyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative, disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mcswiggen J, Morrissey D, Pavco P,
                                                                                                                                     HCV minus strand DNAzyme substrate sequence #801
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Page 289; 387pp; English.
                              ACD62938 standard; RNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-MAR-2001; 2001US-00817879.
08-UUN-2001; 2001US-0087478.
08-UUN-2001; 2001US-0296876P.
24-OCT-2001; 2001US-0335059P.
05-DEC-2001; 2001US-0337055P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RIBOZYME PHARM INC.
BLATT L.
MACEJAK D.
MCSWIGGEN J.
MORRISSEY D.
                                                                                                                                                                                                                                                                                                                                                                                                                                            26-MAR-2002; 2002WO-US009187
                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Macejak D,
Roberts E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (RIBO-) RIBOZYME PHA
BLATT L.
(MASK), MACBJAK D.
(MCSW), MCSWIGGEN J.
(MORK), MORRISSEY D.
(PAVC), PAVCO P.
(LEEP), LEE P.
(LEEP), DRAPER K.
(ROBE)) ROBERTS E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-229207/22.
                                                                                                                                                                                                                                                                                                                                   Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                        40200281494-A1.
                                                                                                     24-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                        17-OCT-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Draper K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          infection.
                                                                   ACD62938;
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RESULT 664
                  4CD6293
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to murine oligonucleotides (ACC62754-ACC68806), which are associated with tumour suppression, tumour reversion, apoptosis and virus resistance. The oligonucleotides are useful as (1) as probes and primers for detecting, identifying, quantifying and/or amplifying nucleic acid, e.g. as one component of a gene chip, in vitro as (anti) sense reagents; and (2) for production of recombinant polypeptides. The oligonucleotides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.
                                                                                                                                                                                                                                                                                                                                                                       Cytostatic; virucide; neuroprotective; noctropic; neuroleptic; murine; tumour suppression; tumour reversion; apoptosis; virus resistance; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease;
                                                                                                                                                                                                                                                                                                                                     Murine oligonucleotide associated with tumour supression, SEQ ID 5492
                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
2.9%; Score 12.4; DB 1; Length 17;
Best Local Similarity 92.9%; Pred. No. 3.8e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
Query Match 2.9%; Score 12.4; DB 1; Length 17; Best Local Similarity 92.9%; Pred. No. 3.8e+02; Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17 BP; 7 A; 5 C; 3 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 673; 738pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tuijnder M;
                                                                                                                                                                                                                     ACC68245 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (MOLE-) MOLECULAR ENGINES LAB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-SEP-2001; 2001FR-00011979.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17-SEP-2002; 2002WO-IB004210.
                                                                             331 CGGACGACCAGGGC 344
                                                                                                                    3 cceaceaccases 16
                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Telerman A, Amson R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-333167/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                       schizophrenia; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO2003025176-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mus musculus.
                                                                                                                                                                                                                                                                                                01-JUL-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-MAR-2003.
                                                                                                                                                                                                                                                          ACC68245;
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Gaps

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4 CACCACTCAGAGAA 17

RESULT 666 ACC65338

48 CACCACTCAGAGGA 61

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ACC65338 standard; DNA; 17
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ACC65338;

(first entry) 01-JUL-2003

Murine oligonucleotide associated with tumour supression, SEQ ID 2585

Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; murine; tumour suppression; tumour reversion; apoptosis; virus resistance; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; ss.

Mus musculus

WO2003025176-A2.

27-MAR-2003

17-SEP-2002; 2002WO-IB004210.

17-SEP-2001; 2001FR-00011979.

(MOLE-) MOLECULAR ENGINES LAB

Tuijnder Amson R, relerman A,

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WPI; 2003-333167/31.

New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.

Disclosure; Page 333; 738pp; French.

The present invention relates to murine oligonucleotides (ACC62754-ACC68806), which are associated with tumour suppression, tumour reversion, apoptosis and virus resistance. The oligonucleotides are useful as grobes and primers for detecting, identifying are quantifying and/or amplifying nucleic acid, e.g. as one component of a gene chip, in vitro as (anti) sense reagents; and (2) for production of recombinant polypeptides. The oligonucleotides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia 

Sequence 17 BP; 3 A; 8 C; 4 G; 2 T; 0 U; 0 Other;

. Match 2.9%; Score 12.4; DB 1; Length 17; Local Similarity 92.9%; Pred. No. 3.8e+02; les 13; Conservative 0; Mismatches 1; Indels Query Match Best Loca Matches

à

RESULT 667

ACC63151/c ID ACC63151 standard; DNA; 17 BP.

ACC63151;

(first entry) 01-JUL-2003

Murine oligonucleotide associated with tumour supression, SEQ ID 398. 

Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; murine; tumour suppression; tumour reversion; apoptosis; virus resistance; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; ss.

Mus musculus

17-SEP-2002; 2002WO-IB004210. WO2003025176-A2 27-MAR-2003. 

17-SEP-2001; 2001FR-00011979.

(MOLE-) MOLECULAR ENGINES LAB

Ë Tuijnder Amson R, Telerman A,

WPI; 2003-333167/31.

New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.

Disclosure; Page 77; 738pp; French.

The present invention relates to murine oligonucleotides (ACC62754-ACC68806), which are associated with tumour suppression, tumour reversion, apoptossis and virus resistance. The oligonucleotides are useful as (1) as probes and primers for detecting, identifying, quantifying and/or amplifying nucleic acid, e.g. as one component of gene chip; in vitro as (anti) sense reagents; and (2) for production of recombinant polypeptides. The oligonucleotides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia

Sequence 17 BP; 3 A; 3 C; 6 G; 5 T; 0 U; 0 Other;

Gaps ô Query Match 2.9%; Score 12.4; DB 1; Length 17; Best Local Similarity 92.9%; Pred. No. 3.8e+02; Matches 13; Conservative 0; Mismatches 1; Indels

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|561/c | ADB43561 standard; DNA; 17 BP. RESULT 668 ADB43561/ ID ADB4

ADB43561;

Tumour suppression/reversion associated nucleotide #3884.

(revised)
(first entry)

18-DEC-2003 04-DEC-2003

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Gaps

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cytostatic; antiviral; neuroprotective; nootropic; neuroleptic; ss; primer; probe; tumour suppression; tumour reversion; apoptosis; virus resistance; transgenic animals; Alzheimer's disease;;schizophrenia; 

Homo sapiens

WO2003040369-A2

15-MAY-2003.

17-SEP-2001; 2001FR-00011981. 17-SEP-2002; 2002WO-IB004219

(MOLE-) MOLECULAR ENGINES LAB

Tuijnder M;

Amson R,

relerman A,

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WPI; 2003-441574/41.
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New nucleic acid encoding human prostate membrane-specific antigen, useful e.g. for treatment of tumors and viral infection, also related polypeptide and antibodies.

Disclosure; Page 486; 771pp; French

The invention relates to the isolation of 6327 nucleotide sequences, fragments of at least 15 consecutive nucleotides of these nucleotides, a fragments of at least 80% identity, after optimal alignment, with the nucleotides, a sequence that hybridizes under stringent conditions with the nucleotides, or the complement, or corresponding RNA, of the nucleotides. The nucleotides are used as probes or primers for detecting, identifying; quantifying and/or amplifying nucleotides, as in vitro sense and antisense sequences, of nucleotides involved in tumour suppression or reversion, apoptosis and or viral resistance, to produce recombinant polypeptides, and to prepare transgenic animals, as experimental models. The nucleotides (also vectors containing the mand cells containing the vectors), the encoded polypeptides and antibodies (AN) against the polypeptide are useful for prevention and/or treatment of viral infections or diseases characterized by development of tumours or cell degeneration (e.g. Alzhelmer's disease or schizophrenia).

Charlysis of the expression of the nucleotides can be used for diagnosis and/or prognosis of these diseases. The nucleotides and polypeptides can be used to screen for their specific interactive molecules.

Charly useful for treating diseases associated with abnormal expression of the nucleotides

Sequence 17 BP; 9 A; 1 C; 5 G; 2 T; 0 U; 0 Other;

Gaps ő Match 2.9%; Score 12.4; DB 1; Length 17; Local Similarity 92.9%; Pred. No. 3.8e+02; les 13; Conservative 0; Mismatches 1; Indels Query Match Best Loca Matches

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ADB45240 standard; DNA; 17 BP. 18-DEC-2003 diagnosis. ADB45240; RESULT 669 ADB452 

(first entry)

Tumour suppression/reversion associated nucleotide #5563.

cytostatic; antiviral; neuroprotective; nootropic; neuroleptic; ss; primer; probe; tumour suppression; tumour reversion; apoptosis; virus resistance; transgenic animals; Alzheimer's disease; schizophrenia;

Homo sapiens.

WO2003040369-A2

15-MAY-2003.

17-SEP-2002; 2002WO-IB004219. 17-SEP-2001; 2001FR-00011981. (MOLE-) MOLECULAR ENGINES LAB

Tuijnder M; Felerman A, Amson R,

WPI; 2003-441574/41.

New nucleic acid encoding human prostate membrane-specific antigen, useful e.g. for treatment of tumors and viral infection, also related

polypeptide and antibodies.

Disclosure; Page 682; 771pp; French.

The invention relates to the isolation of 6327 nucleotide sequences, fragments of at least 15 consecutive nucleotides of these nucleotides, a sequence having at least 80% identity, after optimal alignment, with the nucleotides, a sequence that hybridizes under stringent conditions with the nucleotides. The complement, or corresponding RNA, of the nucleotides or the complement, or corresponding RNA, of the nucleotides are used as probes or primers for detecting, identifying, quantifying and/or amplifying nucleic acids, as in vitro sense and antifense sequences, of nucleotides involved in tumour recombinant polypeptides, and to prepare transgenic animals, as recombinant polypeptides, and to prepare transgenic animals, as recombinant models: The nucleotides (also vectors containing them and calls containing the vectors), the encoded polypeptides and antibodies (Ab) against the polypeptide are useful for prevention and/or treatment or cell degeneration (e.g. Alzhehmer's disease or schizophrenia).

Analysis of the expression of the nucleotides can be used for diagnosis and bused to screen for their specific interactive molecules. expression of the nucleotides.

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Sequence 17 BP; 5 A; 4 C; 5 G; 3 T; 0 U; 0 Other;

Gaps ö 2.9%; Score 12.4; DB 1; Length 17; 92.9%; Pred. No. 3.8e+02; tive 0; Mismatches 1; Indels Query Match 2.9 Best Local Similarity 92.9 Matches 13; Conservative

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RESULT 670 ADE13461/c

ADE13461 standard; DNA; 17 BP.

ADE13461;

(first entry) 29-JAN-2004 HLA class I allele specific primer #77.

ss; primer; PCR; human; Human Leukocyte Antigen; HLA; genotype.

Homo sapiens.

US2003165884-A1.

04-SEP-2003.

99US-0172768P. 25-APR-2002; 2002US-00133779. 20-DEC-1999; 99US-0172768P. 20-DEC-2000; 2000US-00747391. 

(STEM-) STEMCYTE INC. Chow R, Tonai R;

Identifying class I or II Human Leukocyte Antigen genotypes using hybridization and amplification assays. WPI; 2003-874916/81.

Claim 7; SEQ ID NO 79; 66pp; English.

The invention relates to a method of identifying a class I or II Human Leukocyte Antigen (HLA) genotype of a subject using hybridisation and amplification assay. The method is used for determining the HLA genotype

RESULT 67 AAQ22412/

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Human, PRO; EST; expressed sequence tag; PCR primer; hybridisation;
probe; blood coagulation disorder; cancer; cellular adhesion disorder;
secreted protein; transmembrane protein; ss.
                       Human PRO274 PCR forward primer 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                9805-00812229
9805-0081817P-
9805-0081838P-
9805-0081952P-
9805-0081955P-
9805-0082568P-
9805-0082700P-
9805-0082700P-
                                                                                                                                                                                                                                                                                                                                                                                                    9805-0079728F

9805-0079920P

9805-0079923P

9805-0080107P

9805-0080165P

9805-0080154P

9805-0080327P

9805-0080333P

9805-0080333P

9805-0080333P

9805-0081049P

9805-0081070P
                                                                                                                                                                                                      98US-0077450P
98US-0077632P
98US-0077641P-
98US-0077649P-
98US-0077791P*
                                                                                                                                                                                                                                                                               98US-00040220.
98US-0078886P.
98US-0078910P.
98US-0078936P.
98US-0078939P.
07-DEC-1999 (first entry)
                                                                                                          Homo sapiens
                                                                                                                                   WO9946281-A2
                                                                                                                                                                                 08-MAR-1999;
                                                                                                                                                         16-SEP-1999
                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                              ö
                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This oligonuclectide was used in an example of synthesis of 3'- acridine-tailed oligonuclectide from acridine-CPG. Blockage of the 3' terminal phosphodiester bond improves resistance to nucleases in serum-contg. media. The new synthesis method avoids the derivatisation step of prior art methods and the possible loss and difficult separation. See AAQ22411-Q22415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Solid support synthesis of 3'-tailedoligo-nucleotide(s) via linker gp. provides nuclease resistant prods. opt. with intercalation to improve anti-sense bonding to DNA or RNA strand.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                     Gaps
 of a subject. The present sequence represents a HLA class I allele
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                                                                                                                                                                                                                                                                                               Acridine-CPG; nuclease resistance; controlled pore glass; ss.
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Pred. No. 4.3e+02;
0; Mismatches 1; Indels
                                                             Length 17;
                                                                                     1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 18 BP; 3 A; 8 C; 1 G; 6 T; 0 U; 0 Other;
                                   Sequence 17 BP; 2 A; 6 C; 7 G; 2 T; 0 U; 0 Other;
                                                           Query Match
Best Local Similarity 92.9%; Pred. No. 3.8e+02;
Matches 13; Conservative 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tabone JC;
                                                                                                                                                                                                                                                                          3'-acridine-tailed oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 12; Page 38; 78pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Reed MW, Meyer RB, Petrie CR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAZ33902 standard; DNA; 18 BP.
                                                                                                                                                                                                  AAQ22412 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                   91WO-US006143.
                                                                                                                                                                                                                                                                                                                                                                                                                         90US-00574348.
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Best Local Similarity 92.9%;
Matches 13; Conservative
                                                                                                              373 TCCTGGACCGCGAC 386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         33 TGGGACGAAGATGG 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18 TGTGACGAAGATGG 5
                                                                                                                                    14 rccredaccecec 1
                                                                                                                                                                                                                                                  15-JUL-1992 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                             (MICR-) MICROPROBE CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1992-096825/12.
                 specific primer.
                                                                                                                                                                                                                                                                                                                                                                                                                         28-AUG-1990;
10-JUN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                  28-AUG-1991;
                                                                                                                                                                                                                                                                                                                                                                           05-MAR-1992.
                                                                                                                                                                                                                                                                                                                                                  VO9203464-A
                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ33902
                                                                                                                                                                                                                            AAQ22412;
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RESULT 672 AAZ33902/c ID AAZ3390 XX AC AAZ3390

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Human, Ship-2; antisense oligonucleotide; phosphorothioate; detection; inhibition; SH2-containing phosphatidylinositol phosphatase-2; ss.

.18
 /*tag= a
 /note= "phosphorothioate linkages"

Location/Qualifiers

Key modified_base Homo sapiens.

Human Ship-2 phosphorothioate antisense oligonucleotide #30735.

22-MAY-2000 (first entry)

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The present invention describes secreted and transmembrane polypeptides and their polymucleotides. The nucleotide sequences are useful as sources of probes, primers, for chromosome mapping, and for generation of antisense sequences. They can also be used to create transgenic animals. The proteins can be used to treat a variety of diseases and disorders, depending on their function. Diseases that may be treated include blood coagulation disorders, cancers and cellular adhesion disorders. They may also be used to raise antibodies. AAZ33891 to AAZ34338, and AAX41665 to AAX41774 represent polynucleotide and polypeptide sequence given in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New secreted and transmembrane polypeptides and their polymuclectides, useful for treating blood coagulation disorders, cancers and cellular adhesion disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Goddard A, Gurney A, Yuan J, Baker KP, Chen J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Match 2.9%; Score 12.4; DB 1; Length 18; Local Similarity 92.9%; Pred. No. 4.3e+02; les 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 18 BP; 2 A; 8 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 4; Page 183; 530pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                          98US-0087106P.
98US-0087208P.
98US-0094651P.
                                         98US-0084441P

98US-0084630P

98US-0084637P

98US-0084643P

98US-0084643P

98US-0084643P

98US-0084643P

98US-0085433P

98US-0085339P

98US-0085339P

98US-008539P

98US-008582P

98US-008643P

98US-008643P

98US-008643P

98US-008643P

98US-008643P
                                                                                                                                                                                                                                                                                                                                                                                                                                                        98US-0100038P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 215 GAACTCGGTGGCGG 228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18 GAACTCCGTGGCGG 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1999~551358/46.
                                                                    07-MAY-1998;
07-MAY-1998;
07-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-SEP-1998;
                                                                                                                07-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                     22-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Wood WI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                              13-MAY-1
13-MAY-1
                                                                                                                                                                                                                                                                                5-MAY-:
                                                                                                                                                                                                                                                                                                                          8-MAY-
                                                                                                                                                                                                                                     LS-MAY-
                                                                                                                                                                                                                                                                                                           5-MAY
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Antisense oligonucleotides, useful for inhibiting human Ship-2 expression and for detecting nucleic acids encoding Ship-2.

99US-00339964. 99US-00339964.

25-JUN-1999; 25-JUN-1999;

US6025198-A.

15-FEB-2000.

COWBERT LM;

Bennett CF,

WPI; 2000-181819/16.

(ISIS-) ISIS PHARM INC.

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The present invention describes phosphorothioate antisense cligonucleotides that specifically hybridise with, and inhibit the expression of, nucleic acids encoding human Ship-2 (also called SH2-containing phosphatidylinositol phosphatase-2). Also described is a method of inhibiting the expression of Ship-2 in human cells or tissues in vitro comprising contacting the cells with the phosphorothioate antisense oligonuclectides. The phosphorothioate antisense oligonuclectides. The phosphorothioate antisense oligonuclectides to be used to treat animals (especially humans) suspected of having or being prone to a disease or condition associated with Ship-2 expression. The present sequence represents a phosphorothioate antisense oligonucleotide for human Ship-2, from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human biallelic marker upstream amplification primer SEQ ID NO:4482.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human genome; biallelic marker; high density disequilibrium map;
genomic map; haplotype; phenotype; polymorphic base; genotyping;
haplotyping; hybridisation; identification; characterisation;
amplification; single nucleotide polymorphism; SNP; PCR primer;
diagnosis; se.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 2.9%; Score 12.4; DB 1; Length 18; Best Local Similarity 92.9%; Pred. No. 4.3e+02; Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18 BP; 7 A; 4 C; 7 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 3; Col 40; 34pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAZ70126 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          273 GAGCAGGGCGCAC 286
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Gaps ö

AAZ91453 standard; DNA; 18 BP.

RESULT 673 AAZ91453

AAZ91453;

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AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ7740 represent amplification primers for the biallelic markers. The biallelic markers of the invention primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex associations studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical afficacious responses to and side effects from pharmaceutical agents acting on a disease as well as other treatment. Pharmaceutical agents acting on a disease as well as other treatment.

When the second actually given a sequence in the Sequence Listing from the present invention
                                                                                                                                                                                                                                                                                       Novel biallelic markers used to construct a high density disequilibrium map of the human genome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human biallelic marker downstream amplification primer SEQ 1D NO:10930.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches 1; Indels 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human genome, biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyphing; haplotyphing; hybridisation; identification; characterisation; amplification; single nucleotide polymorphism; SNP; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.9%; Score 12.4; DB 1; Length 18; 92.9%; Pred. No. 4.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18 BP; 1 A; 7 C; 1 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                      Chumakov I;
                                                                                                                                                                                                                                                                                                                                           Claim 8; Page 1186; 2745pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAZ76574 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99WO-IB000822.
                                                                                                  99WO-IB000822.
                                                                                                                                    98US-0082614P
                                                                                                                                                   98US-0109732P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             362 CTTCCTCACTTTCC 375
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Matches 13; Conservative
                                                                                                                                                                                                                      Cohen D, Blumenfeld M,
                                                                                                                                                                                                                                                       WPI; 2000-013267/01
                                                                                                                                                                                    GEST ) GENSET,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              diagnosis; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-APR-1999;
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                                                                                                21-APR-1999;
                                                                                                                                    21-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           W09954500-A2
Homo sapiens
                               WO9954500-A2
                                                                                                                                                   23-NOV-1998;
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                                                                 28-OCT-1999.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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AAZ76574/c
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invention, which contains a polymorphic base at position 24 of their nucleotide sequences. AAZ68579 to AAZ7740 represent amplification primers for the biallelic markers of the biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyphag studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and dispossic methods, as well as the characterisation of the pharmaceutical as differential effications responses to and side effects from pharmaceutical agents acting on a disease as well as other treatment.

N.B. The SEQ ID NOS 2852, 2913, 2974, 3055, 3056, 3157, 3227, 3297 and and also actually given a sequence in the Sequence Listing from the
                                                                                                                                               Novel biallelic markers used to construct a high density disequilibrium
                                                                                                                                                                                                                                AAZ65654 to AAZ69578 represent human biallelic markers from the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human, secreted protein, transmembrane protein, PRO, EST, cytostatic, expressed sequence tag, detection, cancer, PCR primer, probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 2.9%; Score 12.4; DB 1; Length 18; Best Local Similarity 92.9%; Pred. No. 4.3e+02; Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 18 BP; 2 A; 7 C; 2 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human PRO274 forward PCR primer SEQ ID NO:14.
                                                                                Cohen D, Blumenfeld M, Chumakov I;
                                                                                                                                                                                                 Claim 9; Page 2561; 2745pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99WO-US005028.
99US-0123957P.
99US-0126773P.
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99US-0131445P.
99US-0134287P.
99US-0145698P.
99US-0145698P.
99US-0162506P.
99WO-US028313.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18-FEB-2000; 2000WO-US004341.
98US-0082614P.
98US-0109732P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 212 AGAGAACTCGGTGG 225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 08-FEB-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15 AGAGAACACGGTGG 2
                                                                                                                                                                   map of the human genome.
                                                                                                               WPI; 2000-013267/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200053756-A2.
                                              (GEST ) GENSET.
21-APR-1998;
23-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-SEP-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAC78608;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 676
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAC78608/c
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The present invention describes a method for distinguishing a human leukocyte antigen (HLA) class I antigen or allele by a combination of bolymerase chain reaction (PCR) using a primer pair whereby all HLA-A, -B or -C alleles can be amplified or using reverse hybridisation analysis comprising a DNA probe covalently bonded to microtitre plate wells which are hybridisable specifically with the base sequence of at least one specific HLA-A, -B or -C allele. The method is applicable in gene typing, judging donor-recipient compatibility during organ transplant and correlation analysis for diagnosis of various diseases. The method is simple, rapid and accurate, with possibility of mechanisation and automation, without the problems encountered by using the prior-art techniques. AAAA66943 to AAAA67072 represent oligonucleotide probes and PCR primers for use in the method of the present invention
                                                                                   Simple, rapid and accurate method for distinguishing HLA class I allele type with possibility of mechanization and automation, applicable in judging donor-recipient compatibility during organ transplant and disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Genetic clustering by distributing members into optimal numbers of clusters determined by a hierarchical clustering algorithm or by paired-pair analysis of homozygous pairs in clusters got from non-hierarchical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cluster; hierarchical clustering algorithm; population based study; clinical trial; DNA fingerprint; genetic profile analysis; PCR primer; SNP; single nucleotide polymorphism; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sample member clustering methed related human DNA PCR primer #28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.9%; Score 12.4; DB 1; Length 18; 92.9%; Pred. No. 4.3e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 18 BP; 5 A; 5 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 61; Page 80; 100pp; English
                                                                                                                                                                                                              Claim 8; Page 66; 83pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        멾
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-OCT-1999; 99US-0161231P.
07-JUL-2000; 2000US-0216897P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAF89291 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         7 GAGTGAAACTGCGG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ä
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Schork N, Skierczynski
  Kaneshige T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-316248/33.
                                          WPI; 2000-400097/34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200129257-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-DEC-2001
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                                                                                                                                                                   diagnosis.
  Moribe T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAF89291;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Loca
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 678
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAC78458 to AAC78599 represent polynucleotide and EST (expressed sequence tag) sequences which encode secreted or transmembrane PRO polypeptides. The PRO polynucleotides and polypeptides have cytostatic activity. The polynucleotides and polypeptides can be used for detecting the presence of PRO polypeptides can be used for detecting the presence of PRO polypeptides on the presence of PRO polypeptides and for modilating bloodstal activities of cells, using the polypeptides for specific targeting. The polypeptide targeting can be used to kill the target cells, e.g. for the treatment of cancers. The polypeptide pairs provide specific targeting of bloactive molecules to cells. AAC78670 to AAC78897 represent PCR primers and probes used in the isolation of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                     Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL;
Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME;
Goddard A, Godowski PJ, Grimaldi CJ, Gurney AL, Hillan KJ;
Kljavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton DL;
Stewart TA, Tumas D, Williams PM, Wood WI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human leukocyte antigen; HLA; class I allele type; probe; PCR primer; amplification; hybridisation; organ transplant; gene typing; dlagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                           Novel PRO polypeptides and polynuclectides used in detection methods, trarget bioactive molecules to specific cells, and to modulate cellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human leukocyte antigen C allele DNA probe B-1 SEQ ID NO:74.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Match 2.9%; Score 12.4; DB 1; Length 18; Local Similarity 92.9%; Pred. No. 4.3e+02; les 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 18 BP; 2 A; 8 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 4; Page 235; 636pp; English.
02-DEC-1999; 99WO-US028565.
16-DEC-1999; 99WO-US0310095.
30-DEC-1999; 99WO-US031243.
30-DEC-1999; 99WO-US031274.
65-JAN-2000; 200WO-US000277.
06-JAN-2000; 200WO-US000377.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAA67016 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98JP-00335151.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99WO-JP005527.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           215 GAACTCGGTGGCGG 228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-OCT-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GAACTCCGTGGCGG 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (SHIO ) SHIONOGI & CO LTD
                                                                                                                                                                                             (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2000-611443/58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200031295-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               32-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     activities.
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RESULT 677 AAA67016

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Gaps

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The present invention relates to a method of determining polynucleotide expression, which comprises hybridising digested cDNA to a capture probe coupled to a solid particle under stringent conditions, where the capture probe is specific for the target polynucleotide and the particle identifies the capture probe. The method is useful for expression profiling, where the presence and/or the amount of a target polynucleotide is similtraneously determined, for diagnosing a disease, condition, disorder, or predisposition associated with a change in expression patterns, in determining the developmental or physiological state of a cell or tissue, for detecting SNPs, which may be used to screen individuals for a genetic predisposition to a disease, condition, or disorder, and in marker assisted selection. The present sequence is a hybridisation tag described in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Determining polymucleotide expression, useful for expressing profiling or detecting single nucleotide polymorphisms, comprises hybridizing digested cDNA, to a capture probe coupled to a solid particle under stringent
            The present invention describes methods of clustering members of a sample, involving applying a hierarchical clustering algorithm to the sample members, determining the optimal number of clusters based on this and distributing the sample members into clusters using non-hierarchical cultstring. The methods are useful in population based studies such as clinical trials, DNA tingerprinting and genetic profile analyses. The present sequence was used to demonstrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleic acid analysis, microarray, single nucleotide polymorphism, SNP; multiplex, expression analysis; hybridisation tag; ss.
                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Drosophila ubx gene SNP analysis universal hybridisation tag #31.
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                                                                                                                                                                                                                   Length 18;
                                                                                                                                                                                                                                                         1; Indels
                                                                                                                                                                           Sequence 18 BP; 3 A; 4 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                 2.9%; Score 12.4; DB 1;
92.9%; Pred. No. 4.3e+02;
tive 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (SYGN ) SYNGENTA PARTICIPATIONS AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hinkel CA, Kimmerly WJ, Yang L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 34; Page 29; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-JAN-2001; 2001US-0264972P. 02-FEB-2001; 2001US-0266186P. 04-JUN-2001; 2001US-0295986P.
                                                                                                                                                                                                                                                                                                                                                                                                                                         AAL49057 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-JAN-2002; 2002WO-EP000868
                                                                                                                                                                                                                                                                                             265 TGCACCTGGAGCAG 278
                                                                                                                                                                                                                                                                                                                                     recacerrasecae 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                         Query Match
Best Local Similarity 92.9
Matches 13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Drosophila sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-OCT-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAL49057;
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Sequence 18 BP; 6 A; 3 C; 5 G; 4 T; 0 U; 0 Other;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             treating
                                                                                                                                                                                                                                              Human, PRO; benign tumour, malignant tumour; lymphoid malignancy, leukaemia, neuronal disorder; stromal disorder; blastocoelic disorder; inflammatory disorder; immune disorder; angiogenic disorder; cytostatic; neuroprotective; PCR; primer; ss.
                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Thirty five nucleic acids encoding PRO polypeptides, useful for benign or malignant tumors, leukemias and lymphoid malignancies, inflammatory, angiogenic and immunologic disorders.
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Stone DM;
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Length 18;
                          Indels
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Smith V,
 Score 12.4; DB 1;
Pred. No. 4.3e+02;
0; Mismatches 1;
                                                                                                                                                                                                                        Forward PCR primer 4 for human PRO274 DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Godowski PJ,
i RM, Roy MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 10; Page 119; 302pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Goddard A, Godowe
Pan J, Pitti RM,
Wood WI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          99WO-US012252.
99US-0140650P.
99US-0140653P.
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99US-0145698P.
99US-0146222P.
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99WO-US028301.
99WO-US028634.
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99US-0133459P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-JAN-2000; 2000WO-US000219
  2.9%;
                                                    230 CAAATCGGGAGGCT 243
                                                                                                                                              ABK40318 standard; DNA; 18
                                                                             CAAAACGGGAGGCT 18
                                                                                                                                                                                                (first entry)
 Query Match
Best Local Similarity 92.9
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99WO-US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-205567/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ashkenazi AJ,
Marsters SA, P
Watanabe CK, W
                                                                                                                                                                                                                                                                                                                                           WO200153486-A1.
                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                              11-FEB-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                               11-MAY-1999;
02-JUN-1999;
22-JUN-1999;
                                                                                                                                                                                                15-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                     26-JUL-2001
                                                                                                                                                                       ABK40318;
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                                                                                                                     RESULT 680
                                                                                                                                  ABK40318/c
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18 AAGCTGCTGAAGGA 5

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The present invention describes a method for determining if an individual has a tumour cell or site of angiogenesis, or if a treatment is effective in changing angiogenesis or changing a status of a set of target cells, comprising determining it sample the subject has an expression product of at least one marker gene. Also described is a compound capable of altering the expression or activity of Keratin 14, TIE 1, Salioadhesin or Siglec, and kits containing them from the present invention can be used in a diagnostic method, particularly as an indicator of angiogenesis or to determine presence of a tumour cell. The method of the invention is suitable to determine within a few days if a certain treatment against Kaposi's Sarcoma is successful. ABQ81851 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Determining presence of a tumor cell or angiogenesis, and the offertiveness of marker genes is
sequences are also useful in gene therapy. The present sequence represents a PCR primer used in the methods of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                 primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            effectiveness of treatment, by detecting the presence of mar)
useful to detect and monitor treatment of Karposl's Sarcoma.
                                                                                 Length 18
                                                                                                                    1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                               Human; Kaposi's sarcoma; tumour; angiogenesis; PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 18 BP; 2 A; 8 C; 3 G; 5 T; 0 U; 0 Other;
                                                  Sequence 18 BP; 2 A; 8 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                 2.9%; Score 12.4; DB 1; 92.9%; Pred. No. 4.3e+02; iive 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                             Kaposi's Sarcoma TAG PCR primer SEQ ID NO:142.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (AMST-) AMSTERDAM SUPPORT DIAGNOSTICS BV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 10; Page 24; 38pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cornelissen M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-JAN-2001; 2001EP-00200228.
28-SEP-2001; 2001EP-00203703.
28-SEP-2001; 2001US-0325722P.
                                                                                                                                                                                                                                                                           ABQ81992 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-JAN-2002; 2002EP-00075264.
                                                                                                                                                      215 GAACTCGGTGGCGG 228
                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                            18 GAACTCCGTGGCGG 5
                                                                              2.99
Best Local Similarity 92.99
Matches 13, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Jan Der Kuyl AC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EP1225233-A2.
                                                                                                                                                                                                                                                                                                                                              19-NOV-2002
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                                                                                                                                                                                                                                                                                                           AB081992;
                                                                                                                                                                                                                                        RESULT 68
ABQ81992/
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first active agent comprising an oligonucleotide antisense to the first active agent comprising an oligonucleotide antisense to the first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genemic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 mucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antisthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing levels of adenosine receptor, producing bronchdilation, increasing levels of adenosine receptor, producing bronchdilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchconscription, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obsained in electronic format directly from WIPO Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or Human, antisense, lung dysfunction, nasal airway dysfunction, antinflammatory steroid, ubiquinone, antinflammatory; antiallergic, antiasthmatic; hypotensive; immunosuppressive, cytostatic; gene therapy; antisense gene therapy; respiratory; lung, adenosine sensitivity; adenosine receptor; bronchodiation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds. ä Nyce JW, Li Y, Sandrasagra A, Katz E, Pabalan J, Aguilar Miller S, Tang L, Shahabuddin S; 2.9%; Score 12.4; DB 1; Length 18; 92.9%; Pred. No. 4.3e+02; tive 0; Mismatches 1; Indels Sequence 18 BP; 2 A; 6 C; 7 G; 3 T; 0 U; 0 Other; Disclosure; SEQ ID NO 12577; 872pp; English. at ftp.wipo.int/pub/published_pct_sequences Human IL4-R oligonucleotide sequence. BP. 23-APR-2002; 2002WO-US013135. 24-APR-2001; 2001US-0286137P. (EPIG-) EPIGENESIS PHARM INC ABZ97335 standard; DNA; 18 (first entry) Query Match
Best Local Similarity 92.9
Matches 13; Conservative WPI; 2003-229219/22. WO200285308-A2. Homo sapiens 17-OCT-2003 31-OCT-2002. ubiquinone. ABZ97335; RESULT ABZ973 

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Gaps

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282 GGCACCAAGCTGGT 295

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Gaps

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Query Match 2.9%; Score 12.4; DB 1; Length 18; Best Local Similarity 92.9%; Pred. No. 4.3e+02; Matches 13; Conservative 0; Mismatches 1; Indels

288 AAGCTGGTGAAGGA 301

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98US-0082804P.
98US-0083132P.
98US-0083132P.
98US-0083132P.
98US-0083132P.
98US-0083149EP.
98US-0083149EP.
98US-0083149EP.
98US-0083149EP.
98US-0083149EP.
98US-0083149EP.
98US-0083149EP.
98US-008313P.
98US-008414P.
98US-008519P.
98US-008519P.
98US-008519P.
98US-008519P.
98US-008518P.
98US-008519P.
98US-008519P.
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98US-008519P.
98US-008519P.
98US-008519P.
98US-008518P.
98US-008518P.
98US-008518P.
98US-0016897B.
98US-0016897B.
98US-0016897B.
98US-0016897B.
98US-0016897B.
98US-0018736B.
98US-0018736B.
99US-0018736B.
99US-0018737B.
99US-0018737P.
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99US-00311832.
99US-0134287P.
   Human; secreted and transmembrane protein; PRO; virucide; gene therapy; cell death; growth induction cascade; blood coagulation cascade; viral infection; PCR; primer; ss.
                                                                                                                                                               Novel human secreted and transmembrane protein related primer #6.
                                                                                                                                                                                                                                                                                                                                                                         9710S-0064249P.
9710S-0064344P.
9710S-0065344P.
9710S-0065344P.
9810S-0077641P.
9810S-0077641P.
9810S-0077641P.
9810S-0077641P.
9810S-0077801P.
9810S-007801P.
9810S-007801P.
9810S-007801P.
9810S-007801P.
9810S-0079664P.
9810S-0079664P.
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9810S-0079664P.
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9810S-007966P.
9810S-0079728P.
9810S-0081077P.
9810S-0081077P.
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9810S-0081077P.
9810S-0081077P.
9810S-0081207P.
9810S-0081207P.
9810S-0081207P.
9810S-0081207P.
9810S-0081207P.
9810S-0081207P.
9810S-0081207P.
                                       RESULT 683
ACD42435 Standard; DNA; 18 BP.
XX
XX
XX
C 90-SEP-2003 (first entry)
XX
W Human; secreted and transmemb:
XX W Human; secreted and transmemb:
XX W Human; secreted and transmemb:
XX W Human; secreted and transmemb:
XX Homo sapiens.
XX USO003050239-Al.
XX USO00305029-Al.
XX USO003050239-Al.
XX USO00303039-Al.
XX USO003050239-Al.
XX USO0030303-Al.
XX U
                                                                                                                                                                                                                                                                                                                                               15-OCT-2001; 2001US-00978191
   1 GGCACCAGGCTGGT 14
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Human; secreted and transmembrane protein; PRO; antiinflammatory; antiarteriosclerotic; cardiant; anti-infertility; anti-HIV; cytostatic; antidiabetic; gene therapy; inflammatory disease; organ failure; atherosclerosis; cardiac injury; infertility; birth defect; premature aging; AIDS; cancer; diabetic complication; chromosome mapping; gene mapping; pharmaceutical; diabetic; biosensor; bioreactor; tissue typing; PCR; primer; ss.
                                                                              Novel human secreted and transmembrane protein related primer #6
                                                                                                                                                                                                                                                                                                                                                                                                            970S-0064249P
970S-006424PP
970S-0066364PP
980S-0077450P
980S-0077641PP
980S-0077641PP
980S-0077641PP
980S-0077641PP
980S-0077641PP
980S-0078034PP
980S-0078034PP
980S-0079664PP
980S-0080334PP
980S-008034PP
980S-0081819PP
980S-0081819PP
980S-0081819PP
980S-0081819PP
980S-0081819PP
980S-0081819PP
980S-0081819PP
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                                         16-JUN-2003 (first entry)
                                                                                                                                                                                                                                                                                                  US2002192706-A1.
                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                        19-DEC-2002
       ACA63470;
       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Baton DL;
Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 2.9%; Score 12.4; DB 1; Length 18; Best Local Similarity 92.9%; Pred. No. 4.3e+02; Matches 13; Conservative 0; Mismatches 1; Indels
14-MAY-1999; 99WO-USO10733.
16-JUN-1999; 99WO-USO10733.
23-JUN-1999; 99US-01367FP.
23-JUN-1999; 99US-01367FP.
26-JUL-1999; 99US-014569BP.
26-JUL-1999; 99US-0146522P.
28-JUL-1999; 99US-0146528P.
25-AUG-1999; 99US-0146522P.
25-AUG-1999; 99US-01380142.
25-AUG-1999; 99US-01380142.
25-AUG-1999; 99US-01380142.
25-AUG-1999; 99US-01380142.
25-AUG-1999; 99US-01380142.
30-NOV-1999; 99US-01380142.
30-NOV-1999; 99US-01380142.
30-NOV-1999; 99WO-USO1855.
16-DEC-1999; 99WO-USO1856.
11-FEB-2000; 2000WO-USO1856.
11-FEB-2000; 2000WO-USO1841.
10-MAX-2000; 2000WO-USO1841.
11-MAX-2000; 2000WO-USO1864.
11-MAX-2000; 2000WO-USO1864.
11-MAX-2000; 2000WO-USO1864.
11-MAX-2000; 2000WO-USO1864.
11-MAX-2000; 2000WO-USO186.
11-MAX-2000; 2000WO-USO186.
11-MAX-2001; 2001WO-USO186.
11-MAX-2001; 2001WO-USO186.
11-MAX-2001; 2001WO-USO186.
11-MAX-2001; 2001WO-USO186.
11-MAX-2001; 2001WO-USO196.
11-MAX-2001; 2001WO-U
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Gaps ö

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The invention describes an isolated nucleic acid (I) comprising, or which is at least 80 % sequence identity to, or the full-length coding sequence off, any of 118 300-2100 nucleotide sequences, which encodes its corresponding PRO polypeptide selected from 118 100-700 amino acid sequences, all given in the specification. The nucleic acids and polypeptides are useful for treating inflammatory diseases, organ failure, atherosclerosis, cardiac injury, infertility, birth defects, premature aging, AIDS, cancer, or diabetic complications. The nucleic acids are useful as hybridisation probes, in chromosome and gene mapping, and in generating antiseanse RNA or NNA. The polypeptides are useful as pharmaceuticals, diagnostics, blosensors or bioreactors. Both are useful in tissue typing. This sequence represents a novel human secreted and transmembrane PRO polypeptide associated primer
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Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME;
Goddard A, Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ;
Kljavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton DL;
Stewart TA, Tumas D, Williams PM, Wood WI;
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20-NOV-1998; 98WO-USO24855.

08-JAN-1999; 99WO-USO0106.

08-JAN-1999; 99WO-USO0106.

10-MAR-1999; 99WO-USO0129.

20-JUN-1999; 99WO-USO1252.

30-NOV-1999; 99WO-USO1252.

30-DEC-1999; 99WO-USO1252.

16-DEC-1999; 99WO-USO1855.

11-FEB-2000; 2000WO-USO1856.

11-FEB-2000; 2000WO-USO1855.

21-MAR-2000; 2000WO-USO1853.

21-MAR-2000; 2000WO-USO1891.

22-MAY-2000; 2000WO-USO1891.

23-MAY-2000; 2000WO-USO1855.

24-AUG-2000; 2000WO-USO1855.
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2001WO-US017092.
2001WO-US017800.
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09-JUL-2001; 2001WO-US021735
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25-MAY-2001
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Score 12.4; DB 1; Length 18; Pred. No. 4.3e+02;

2.9%;

Query Match Best Local Similarity

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Human; ds; thrombolytic agent; interferon; interleukin; cytokine; erythropoietin; colony stimulating factor; cancer; colorectal carcinoma; appprosis related condition; AIDS; amyotrophic lateral sclerosis; inflammatory disease; asthma; atherosclerosis; neurodegenerative disease; gastrointestinal disorder; Alzheimer's disease; Parkinson's disease; paperrointestinal disorder; Alzheimer's disease; Parkinson's disease; plypertension; myocardial ischaemia; Kidney disease; carcinogenesis; glomerulonephritis; lung disease; pulmonary hypertension; preeclampsia; bronchial asthma; gastric ulcer; renal failure; cardiovascular disease; inflammatory bowel disease; reproductive disorder; premature labour.
                                                                                                                                                                  Human PRO polypeptide associated oligonucleotide SEQ ID NO 14.
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Mismatches
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97US - 0065311P
97US - 0065311P
98US - 0077450P
98US - 0077641P
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9805-0078910P.
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98US-00105413.
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98US-00184216.
98US-00187368.
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98US-00202054.
98US-00218517.
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                      215 GAACTCGGTGGCGG
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26-JUN-1998;
07-OCT-1998;
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27-MAR-19
27-MAR-19
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ACA71634/
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New isolated PRO polypeptides e.g. PRO213, PRO274 and PRO300, for use as pharmaceuticals, diagnostics, biosensors and bioreactors, for identifying modulators of receptor-ligand interactions.
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Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME;
Goddard A, Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ;
Kljavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton DL;
Stewart TA, Tumas D, Williams PM, Wood WI;
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12-APR-1999; 99US-00284291.
14-MAY-1999; 99WG-00311832.
12-MAY-1999; 99WG-00311832.
22-AUG-1999; 99US-00380137.
25-AUG-1999; 99US-00380137.
25-AUG-1999; 99US-00380137.
25-AUG-1999; 99WG-US028313.
30-NOV-1999; 99WG-US028513.
30-DEC-1999; 99WG-US028565.
16-DEC-1999; 99WG-US028565.
30-DEC-1999; 99WG-US030095.
30-DEC-1999; 99WG-US030095.
30-DEC-1999; 99WG-US030095.
30-DEC-1999; 99WG-US030095.
30-DEC-1999; 99WG-US030095.
30-DEC-1999; 99WG-US030095.
31-DEC-1999; 99WG-US030095.
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28-JUL-2000; 2000MO-USG15264.
28-JUL-2000; 2000MO-USG22310.
28-JUL-2000; 2000MO-USG23312.
08-NOV-2000; 2000WC-USG23312.
01-DEC-2000; 2000WC-USG3499.
01-DEC-2000; 2000WC-USG3499.
20-DEC-2000; 2000WC-USG3499.
22-MAR-2001; 2001WS-00816744.
22-MAR-2001; 2001WS-0081674.
22-MAR-2001; 2001WS-USG16952.
23-MAR-2001; 2001WS-USG16952.
23-MAR-2001; 2001WS-USG16952.
24-MAR-2001; 2001WS-USG16953.
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21-MAR-2000; 2000WO-US007532.
30-MAR-2000; 2000WO-US008439.
17-MAY-2000; 2000WO-US013705.
22-MAY-2000; 2000WO-US014042.
30-MAY-2000; 2000WO-US014941.
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05-JUN-2001; 2001US-00874503
14-JUN-2001; 2001US-00882636
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30-JUL-2001; 2001US-00918585
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Eaton DL; , Gerritsen ME;

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pharmaceuticals, diagnostics, biosensors and bioreactors. The PRO polypeptide is also useful as a thrombolytic agent, interferon, chiraleukin, erythropoletin, colony stimulating factor and other cytochines. The PRO polypeptide is useful for treating disease such as crocer e.g. colorectal carcinoma; apoptosis related conditions e.g. AIDS, cancer e.g. colorectal carcinoma; inflammatory disease e.g. asthma, carborosclerosis; neurodegenerative disease e.g. Alzheimer's disease, Parkinson's disease; cardiovascular disease e.g. hypertension and glomerulonephritis; lung disease e.g. pulmonary hypertension and glomerulonephritis; lung disease e.g. pulmonary hypertension, bronchial asthma; gastrointestinal disorders e.g. pulmonary hypertension, bronchial component disease; reproductive disorders e.g. pulmonary hypertension, bronchial asthma; carcinogenesis. The present sequence represents a PRO polypeptide associated oligomucleotide of the invention. Note: The sequence data for this patent did not form part of the printed sequence atta for this patent did not form part of the printed special and the present control format directly from USPTO cat sequence. html?DocID=20020177553
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970S-0062250P.
970S-0065311P.
980S-0077632P.
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les 13, Conservative
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11-MAR-1998;
11-MAR-1998;
11-MAR-1998;
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17-OCT-1997;
03-NOV-1997;
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Matches 13;
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08-NOV-2000; 2000US-00709238.
27-NOV-2000; 2000US-0072749.
01-DEC-2000; 2000WG-US032678.
20-DEC-2000; 2000WG-00747259.
20-DEC-2000; 2000WG-09034956.
28-FEB-2001; 2001WG-US005520.
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2001US-00886342.
2001WO-US019692.
2001WO-US021066.
2001WS-US021735.
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2001WO-US017800.
2001US-00874503.
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99WG-US00508.
99WG-US005190.
99WG-US005190.
99WG-US005190.
99WG-US010133.
99WG-US010133.
99WG-US010133.
99WG-US010137.
99WG-US010137.
99WG-US010137.
98US-0079664P.
98US-0079664P.
98US-0079689P.
98US-007928P.
98US-0079786P.
98US-0079923P.
98US-0079923P.
                                                                             98WO-USO21141.
98US-00184216.
98US-00187368.
98WO-USO24855.
98US-00202054.
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2000WO-US020710.
2000WO-US023328.
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22-MAR-2001; 2001US-00816920.
22-MAR-2001; 2001WO-US009552.
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27-MAR-1998,
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31-MAR-1999,
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31-MAR-1999,
32-MAR-1999,
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(GETH ) GENENTECH INC

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The present invention relates to the isolation of novel human PRO CO polypeptides are secreted and transmembrane proteins. The PRO polypeptides are secreted and transmembrane proteins. The PRO polypeptides are useful for detecting other PRO polypeptides, for linking to polypeptides are useful for detecting other PRO polypeptides, for modulating bloogical activities of cells expressing PRO polypeptides, and for for identifying agonists or antagonists. The bloactive molecule maybe a coxin, radiolabel or antibody, and causes apoptosis or death of the cell. The PRO polypeptides are useful for treating immune disorders, diabetes or hypo-insulinaemia, cardiac insufficiency, nervous system cisorders, kidney disorders, bone and cartilage disorders or arthritis, tumours, and wound healing. The polymoclectide sequences encoding PRO polypeptides are useful as hybridiation probes, in chromosome and gene compaping, in the generation of antisense RNA and DNA, in the preparation of propagation of antisense RNA and DNA, in the preparation of an immals, for the generation of antisense RNA and DNA, in the preparation of an immals, for the generatic sequence represents a PCR primer used in the examples of the present invention. Note: The sequence data for this compactual as equent as obtained in electronic format directly from the USPTO web site are sequence.
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Ashkenazi A, Baker KP, Botstein D, Desnoyers L, Eaton D;
Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritssen ME;
Goddard A, Goddowski PJ, Grimaldi JC, Gurney AL, Hillan KJ;
Kljavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton Stewart IA, Tumas D, Williams PM, Wood WI;
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                                                                                                                                                                           Novel secreted and transmembrane polypeptides and polynucleotides encoding them useful for treating cancer, kidney diseases, bone, cartilage disorders and immune deficiencies
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2.9%; Score 12.4; DB 1; Length 18;
Best Local Similarity 92.9%; Pred. No. 4.3e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
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97US-0064249P
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03-NOV-1997;
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PR 13-NOV-1997; 97US-0066361P.
PR 21-NOV-1997; 97US-0066361P.
PR 11-YAR-1998; 98US-0077461P.
PR 11-YAR-1998; 98US-0077641P.
PR 11-YAR-1998; 98US-0077641P.
PR 12-YAR-1998; 98US-0077641P.
PR 20-YAR-1998; 98US-007791P.
PR 20-YAR-1998; 98US-0078936P.
PR 20-YAR-1998; 98US-0078936P.
PR 20-YAR-1998; 98US-0078939P.
PR 20-YAR-1998; 98US-0079868P.
PR 20-YAR-1998; 98US-0079868P.
PR 21-YAR-1998; 98US-0079868P.
PR 21-YAR-1999; 99US-0079861P.
PR 21-YAR-1999; 99US-00798113.
PR 22-YAR-1999; 99US-00380113.
PR 21-YAR-1999; 99US-0038013.
PR 21-YAR-2000; 2000WO-US00331.
PR 21-YAR-2000; 2000
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Query Match 2.9%; Score 12.4; DB 1; Length 18; Best Local Similarity 92.9%; Pred. No. 4.3e+02; Matches 13; Conservative 0; Mismatches 1; Indels

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Gaps

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Human; secreted and transmembrane protein; PRO; tissue typing; chromosome identification; vaccine; cancer; retinal disorder; sports-related joint disorder; osecoarthitis; theumatoid arthritis; wound healing; obesity; diabetes; hearing loss; hearing loss cardiac insufficiency disorder; kidney disorder; nervous system disorder; haemoglobin associated disorder; PCR; primer; ss.
                                                                                                                                                                                                                 Secreted and transmembrane PRO protein associated primer #8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            970S-0064249P
970S-0064249P
970S-0066341P
980S-0077449P
980S-0077649P
980S-0077649P
980S-0077649P
980S-0077649P
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980S-0079668P
980S-0079668P
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980S-0079723P
980S-0080107P
980S-008107P
980S-008107P
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980S-008107P
980S-0081839P
                                                                                                      ADA24553 standard; DNA; 18 BP
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18 GAACTCCGTGGCGG S
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03-NOV-1997;
13-NOV-1997;
13-NOV-1997;
11-MAR-1998;
11-MAR-1998;
11-MAR-1998;
12-MAR-1998;
20-MAR-1998;
20-MAR-1998;
20-MAR-1998;
20-MAR-1998;
27-MAR-1998;
27-MAR-1998;
27-MAR-1998;
27-MAR-1998;
27-MAR-1998;
31-MAR-1998;
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01-APR-1998;
01-APR-1998;
08-APR-1998;
08-APR-1998;
09-APR-1998;
09-APR-1998;
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21-APR-1998;
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                                                                   RESULT 688
ADA24553/c
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PR 22-APR-1998 | 98US-0082704P
PR 22-APR-1998 | 98US-0082704P
PR 22-APR-1998 | 98US-0082704P
PR 22-APR-1998 | 98US-0082704P
PR 23-APR-1998 | 98US-00823135P
PR 29-APR-1998 | 98US-00831322P
PR 29-APR-1998 | 98US-0083132P
PR 29-APR-1998 | 98US-0084649P
PR 07-MAY-1998 | 98US-0084641P
PR 07-MAY-1998 | 98US-0084641P
PR 07-MAY-1998 | 98US-0084641P
PR 13-MAY-1998 | 98US-0084641P
PR 13-MAY-1998 | 98US-0084641P
PR 13-MAY-1998 | 98US-0084641P
PR 13-MAY-1998 | 98US-0084641P
PR 22-MAY-1998 | 98US-008532P
PR 13-MAY-1998 | 98US-008532P
PR 13-MAY-1998 | 98US-008532P
PR 13-MAY-1998 | 98US-008532P
PR 22-MAY-1998 | 98US-008532P
PR 23-MAY-1998 | 98US-008532P
PR 23-MAY-1999 | 98US-010858P
PR 23-MAY
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Human; secreted and transmembrane protein; PRO; cell death; neuropathy; peripheral neuropathy; diabetic peripheral neuropathy; AIDS-associated neuropathy; Charcot-Marie-Tooth disease; Actefueum's disease; Abetalipoproteinaemia; Tangier disease; Krabbe's disease; Metachromatic leukodystrophy; Pabry's disease; Dejerine-Sottas syndrome; chromosome mapping; gene mapping; gene therapy; PCR; primer; ss.
                                                                                       Novel human secreted and transmembrane protein related primer #6.
                                                                                                                                                                                                                                                                                                                                                                                                             9705-0064249P-
9705-00663449P-
9705-00663149P-
9705-0066314P-
9805-0077641P-
9805-0077641P-
9805-0077649P-
9805-0077649P-
9805-0077649P-
9805-0078866P-
9805-0078866P-
9805-0078936P-
9805-00799664P-
9805-0079964P-
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9805-0079938P-
9805-0080337P-
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9805-008123P-
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                                                         (first entry)
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27-WAR-1998;
27-WAR-1998;
27-WAR-1998;
27-WAR-1998;
30-WAR-1998;
31-WAR-1998;
31-WAR-1998;
31-WAR-1998;
31-WAR-1998;
                                                         08-SEP-2003
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                     ACD29616;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New isolated PRO polypeptides for example extracellular, secreted and membrane bound proteins, useful for modulating the biological activities of cells and for treating, for example diabetes, cancer, rheumatoid arthritis, and hearing joss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Shelton DL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention describes an isolated secreted and transmembrane (PRO) polypeptide (I). PRO337 polypeptide is useful for detecting PRO4993 polypeptide in a sample, and vice versa. PRO725, PRO700 and PRO4993 useful for detecting PRO155, PRO700 and PRO159 are useful for detecting PRO155, PRO700 and PRO393 is useful for linking a bioactive molecule to a cell expressing a PRO337 polypeptide, and PRO337 is useful for linking a bioactive molecule to a cell expressing a PRO493 polypeptide. PRO1559 is useful for linking a bioactive molecule for bioactive molecule collecule of a cell expressing a PRO493 polypeptide. PRO1559 is useful for linking a bioactive molecule et all expressing a PRO493 polypeptide. PRO1559 PRO700 and PRO739 polypeptides are useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    KP, Botstein D, Desnoyers L, Eaton DL; E, Fong S, Gao W, Gerber H, Gerritsen ME; PJ, Grimaldi JC, Gurney AL, Hillan KJ; Walber MA, Pan J, Paoni NP, Roy MA, Shelto Williams PM, Wood WI;
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30-NOV-1999; 99WO-USD28313.

02-DEC-1999; 99WO-USD28551.

16-DEC-1999; 99WO-USD28555.

30-DEC-1999; 99WO-USD31243.

30-DEC-1999; 99WO-USD31243.

30-DEC-1999; 99WO-USD31243.

30-DEC-1999; 99WO-USD31243.

30-DEC-1999; 99WO-USD31274.

05-JAN-2000; 2000WO-USD00376.

11-FEB-2000; 2000WO-USD00376.

11-FEB-2000; 2000WO-USD00376.

11-FEB-2000; 2000WO-USD00319.

11-MAR-2000; 2000WO-USD06319.

11-MAR-2000; 2000WO-USD06319.

11-MAR-2000; 2000WO-USD13705.

11-MAR-2000; 2000WO-USD13705.

11-MAR-2000; 2000WO-USD13706.

11-MAR-2000; 2000WO-USD13706.

11-MAY-2000; 2000WO-USD13706.

11-MAR-2000; 2000WO-USD13706.

11-MAR-2000; 2000WO-USD37670.

11-MAR-2000; 2000WO-USD37670.

11-MAR-2000; 2000WO-USD37670.

11-MAR-2001; 2010WO-USD37670.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-JUN-2001; 2001WO-US017800
20-JUN-2001; 2001WO-US019692
29-JUN-2001; 2001WO-US021066
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30-JUL-2001; 2001US-00918585
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      215 GAACTCGGTGGCGG 228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AJ, Baker KP, F, Filvaroff E, I, Godowski PJ, CJ, Kuo SS, Napie A, Tumas D, Will
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Ferrara N, F
Goddard A, G
Kljavin IJ,
Stewart TA,
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RESULT 689 ACD29616/c ID ACD29616 standard; DNA; 18 BP.

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Query Match Local Best Loca Matches

98US-00834999-98US-00835609-98US-00835545-98US-00835589-98US-0083589-98US-0084562-98US-00844149-98US-00844119-98US-008454119-98US-008454119-98US-008454119-98US-008454119-

28-APR-1998;
29-APR-1998;
29-APR-1998;
29-APR-1998;
29-APR-1998;
29-APR-1998;
29-APR-1998;
29-APR-1998;
30-APR-1998;
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30-APR-1998;
30-APR-1998;
30-APR-1998;
30-APR-1998;
30-APR-1998;
31-APR-1998;

15-MAY-1998

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The invention describes an isolated, secreted and transmembrane polypeptide, termed PRO polypeptide (I). (I) is useful for detecting PRO4993, PRO3179, PRO4993, PRO725, PRO7025, PRO7039 polypeptide, and for linking a bioactive molecule to a call expressing the above polypeptides. The bioactive molecule is a toxin, radiolabel or an antibody and causes cell death. (I) is useful as therapeutic agent, in medical and industrial applications e.g. for treating neuropathy, especially peripheral neuropathy, dibS-associated neuropathy, charcot-Marie-Tooth disease, Refusum's disease, Abetalipoproteinaemia, Tangier disease, Krabbe's disease, Metachromatic leukodystrophy, Pabry's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL; Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME; Goddard A, Godowski PJ, Grimaldh JC, Gurney AL, Hillan KJ; Kliavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton DL; Stewart TA, Tumas D, Williams PM, Wood WL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel secreted and transmembrane polypeptide for modulating biological activity of cell expressing the polypeptide, identifying agonists or antagonists of polypeptide, and as molecular weight markers.
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2.9%; Score 12.4; DB 1; Length 18;
Best Local Similarity 92.9%; Pred. No. 4.3e+02;
Matches 13; Conservative 0; Mismatches 1; Indels ...
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                                                                                                                                                                                                                                                                                         28-FEB-2001; 2001WO-US0056520.
22-MAR-2001; 2001WO-US005652.
25-MAR-2001; 2001WO-US01595.
01-UJN-2001; 2001WO-US017800.
20-UJN-2001; 2001WO-US01565.
29-UJN-2001; 2001WO-US021066.
03-UJN-2001; 2001WO-US021735.
30-UJL-2001; 2001US-00918585.
              2000WO-US000277.
2000WO-US000376.
2000WO-US004341.
2000WO-US005804.
2000WO-US005812.
2000WO-US005812.
2000WO-US005812.
2000WO-US005813.
2000WO-US013705.
2000WO-US014042.
2000WO-US012705.
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2000WO-US032678.
2000WO-US034956.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GETH ) GENENTECH INC.
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                              06-JAN-2000; 2
11-FEB-2000; 2
14-FEB-2000; 2
24-FEB-2000; 2
02-MAR-2000; 2
10-MAR-2000; 3
11-MAR-2000; 2
31-MAR-2000; 2
31-MAY-2000; 2
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98US-0084640P 98US-0085338P 98US-0085338P 98US-0085339P 98US-0085533P 98US-0085573P 98US-00855779P 98US-008568P 98US-008568P 98US-008568P 98US-008568P 98US-008568P 98US-008568P 98US-008568P 98US-008648P 98US-008648P 98US-008648P 98US-008709B 98US-01359P 98US-01359P 98US-013631P 98US-013631P 98US-013631P 98US-013631P 98US-013631P 98US-013631P 98US-013631P 98US-013631P 99US-013631P 99US-013631P

12-MAR-1999; 29-MAR-1999; 21-APR-1999;

99WO-US031243 99WO-US031274

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primer; ss; inflammatory disease; organ failure; atherosclerosis; cardiac injury; infertility; birth defect; premature aging; AIDS; cancer; diabetic complication; tissue typing; human; PCR.
                                                                                                        96US-0018049P

97US-0062250P

97US-006434P

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98US-0077454P

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98US-0080334P

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98US-0081819P

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98US-00831845P

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98US-00831845P

98US-00831845P
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                                                                                                        21-MAY-1996;
17-OCT-1997;
13-NOV-1997;
13-NOV-1997;
11-MAR-1998;
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12-MAR-1998;
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33-MAR-1998;
31-MAR-1998;
32-MAR-1998;
                                                                                       17-0CT-2001;
                                   Homo sapiens
                                                                      20-MAR-2003
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PR 30.APR-1998; 98US-0083742P.
PR 06-MAY-1998; 98US-0084366P.
PR 06-MAY-1998; 98US-0084441P.
PR 77-MAY-1998; 98US-0084660P.
PR 77-MAY-1998; 98US-008463P.
PR 77-MAY-1998; 98US-008463P.
PR 77-MAY-1998; 98US-008463P.
PR 77-MAY-1998; 98US-008463P.
PR 13-MAY-1998; 98US-008653P.
PR 13-MAY-1998; 98US-008563P.
PR 22-MAY-1998; 98US-00863P.
PR 22-MAY-1999; 98US-00863P.
PR 22-MAY-1999; 98US-00863P.
PR 22-MAY-1999; 98US-003052P.
PR 22-MAY-1999; 98US-003052P.
PR 22-MAY-1999; 99US-01335P.
PR 22-MAY-1999; 99US-01335P.
PR 23-MAY-1999; 99US-01335P.
PR 23-MAY-1999;

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diabetic peripheral neuropathy; autonomic neuropathy; reduced motility of the gastrointestinal tract; atony of the urinary bladder; post polio syndrome; Krabbe's disease; Charcot-Marie-Tooht disease; Fabry's disease; Tangier disease; Refsum's disease; PCR; primer; ss.
cardiac- insufficiency disorder; peripheral neuropathy;
                                                                                                                                                                                                                                    97US-0064249P.
97US-0065341P.
97US-0065341P.
98US-0077450P.
98US-0077451P.
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98US-0080333P
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9805-0081817P
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9805-0083495P
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29-APR-1998;
29-APR-1998;
                                                                                                              Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Botstein D, Desnoyers L, Eaton DL;
Fong S, Gao W, Gerber H, Gerritsen ME;
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                                                                                                                            18-FEB-2000; 2000MC-0S004341.
24-FEB-2000; 2000MC-0S004341.
10-MAR-2000; 2000MC-0S00541.
11-MAR-2000; 2000MC-0S00541.
11-MAR-2000; 2000MC-0S00541.
11-MAY-2000; 2000MC-0S014042.
11-MAY-2000; 2000MC-0S014042.
11-MAY-2000; 2000MC-0S014042.
12-MAY-2000; 2000MC-0S014042.
13-MAY-2000; 2000MC-0S014042.
14-MC-2000; 2000MC-0S014041.
16-MAR-2000; 2000MC-0S014041.
16-MAR-2000; 2000MC-0S014041.
16-MAY-2001; 2001MC-0S014041.
16-MAY-2001; 2001MC-0S014041.
16-MAY-2001; 2001MC-0S014041.
16-MAY-2001; 2001MC-0S014041.
16-MAY-2001; 2001MC-0S014041.
16-MAY-2001; 2001MC-0S014092.
 02-DEC-1999; 99WO-USO28565.
16-DEC-1999; 99WO-USO31095.
30-DEC-1999; 99WO-USO31243.
30-DEC-1999; 99WO-USO31243.
30-DEC-1999; 99WO-USO31274.
65-JAN-2000; 200WO-USO00277.
66-JAN-2000; 200WO-USO00376.
11-FEB-2000; 200WO-USO0376.
11-FEB-2000; 200WO-USO0376.
11-FEB-2000; 200WO-USO0376.
11-FEB-2000; 200WO-USO0376.
11-FEB-2000; 200WO-USO0376.
11-MAR-2000; 200WO-USO0376.
11-MAR-2000; 200WO-USO0376.
11-MAR-2000; 200WO-USO13705.
11-MAR-2000; 200WO-USO14042.
11-MAR-2000; 200WO-USO13705.
11-MAR-2000; 200WO-USO13705.
12-MAY-2000; 200WO-USO13705.
13-MAY-2000; 200WO-USO13705.
13-MAY-2000; 200WO-USO13705.
14-MAY-2000; 200WO-USO13705.
15-MAY-2000; 200WO-USO13705.
16-MAY-2000; 200WO-USO13705.
17-MAY-2000; 200WO-USO13705.
18-MAY-2000; 200WO-USO13705.
18-MAY-2000; 200WO-USO13705.
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2001US-00882636.
2001US-00886342.
2001WO-US019692.
2001WO-US021066.
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Ferrara N, Filvaroff E,
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Best Local Similarity 92.9
Matches 13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-JUN-2001; 20
05-JUN-2001; 20
14-JUN-2001; 20
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ACD29031/c
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98US-0083554P.
98US-0083558P.
98US-0084558P.
98US-008458P.
98US-0084541P.
98US-0084637P.
98US-0084637P.
98US-0084637P.
98US-0084637P.
98US-0084637P.
98US-0084637P.
98US-0084637P.
98US-00858P.
98US-0085873P.
98US-00858P.
98US-0085873P.
98US-0085823P.
98US-008582P.
98US-0085823P.
98US-0085823P.
98US-0085823P.
98US-0085823P.
98US-0085823P.
98US-0085823P.
98US-0085823P.
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99US-013322P.
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99US-0131322P.
99US-0131322P.
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99US-01313287P.
99US-01313287P.
99US-0131287P.
99US-01313287P.
99US-0131287P.
99US-0131287P.
29-APR-1998;
29-APR-1998;
30-APR-1998;
30-APR-1998;
66-MAY-1998;
66-MAY-1998;
67-MAY-1998;
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01-JUL-1998
11-SEP-1998
07-OCT-1998
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02-NOV-1998
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20-NOV-1998
20-NOV-1998
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Gaps
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2.9%; Score 12.4; DB 1; Length 18;
Best Local Similarity 92.9%; Pred. No. 4.3e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
29-OCT-1999; 99WS-0162506P.

20-NOV-1999; 99WO-US028513.

02-DEC-1999; 99WO-US028551.

02-DEC-1999; 99WO-US028551.

03-DEC-1999; 99WO-US031024.

30-DEC-1999; 99WO-US031024.

30-DEC-1999; 99WO-US031274.

10-MAN-2000; 2000WO-US000277.

11-FEB-2000; 2000WO-US00376.

11-FEB-2000; 2000WO-US00376.

12-MAR-2000; 2000WO-US00376.

10-MAR-2000; 2000WO-US003441.

10-MAR-2000; 2000WO-US00341.

10-MAR-2000; 2000WO-US01441.

10-MAR-2000; 2000WO-US01441.

10-MAR-2000; 2000WO-US01441.

10-MAR-2000; 2000WO-US01441.

10-MAR-2000; 2000WO-US013706.

21-MAR-2000; 2000WO-US013706.

22-MAR-2000; 2000WO-US013706.

24-AUG-2000; 2000WO-US013779.

24-AUG-2000; 2000WO-US013779.

25-MAR-2000; 2000WO-US013779.

26-UN-2000; 2000WO-US013779.

27-NOV-2000; 2000WO-US012779.

27-NOV-2000; 2000WO-US01779.

27-NOV-2000; 2000WO-US01
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AC
ADB73520 standard; DNA; 18 BP
AC
ADB73520;
XX
XX
Human, PRO DNA PCR primer #6.
XX
KW Human, PRO polypeptide; secre
KW Cell death; neuropathy; neuro
KW Call death; neuropathy; ne
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16-OCT-2001; 2001US-00978608

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250P 249P 311P	632P 632P 641P 649P	004P	2397 2397 2397 1656 1656	6637 6647 6897 7287	9207 9237 1057 1077 1657	194P	10495 10705 10715 11955 12035	1817 1819 1952 1955 1955 1955 1955 1955 1955 19	22569E 2700E 2704E 2797E 3336E	88888888888888888888888888888888888888	8846279. 8846379. 8846209. 88464114. 8846209. 8846379. 88464379. 8846439.
US-006 US-006 US-006 US-006	US-007	US-007 US-007 US-007 US-007	US-007 US-007 US-007 US-007	TUS-007 TUS-007 TUS-007 TUS-007	3US-007 3US-008 3US-008 3US-008	3US-008 3US-008 3US-008 3US-008	800-SD8	302-008 303-008 303-008 303-008	300-SD8 300-008 300-008 300-008 300-008	800-508 803-008 803-008 803-008 803-008	800 - S.086 800 - S.086
2-2-2-2											
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PR 13-MAY-1998 98US-0085339P.
PR 15-MAY-1998 98US-0085339P.
PR 15-MAY-1998 98US-0085533P.
PR 15-MAY-1998 98US-0085532P.
PR 15-MAY-1998 98US-0085532P.
PR 15-MAY-1998 98US-0085582P.
PR 15-MAY-1998 98US-0085682P.
PR 15-MAY-1998 98US-0085632P.
PR 22-MAY-1998 98US-0085430P.
PR 22-MAY-1998 98US-001010P.
PR 22-MAY-1998 98US-001010P.
PR 22-MAY-1999 98US-0010139P.
PR 22-DEC-1998 98US-0113328P.
PR 22-DEC-1999 99US-0113328P.
PR 22-DEC-1999 99US-01336128P.
PR 22-DEC-1999 99US-01336128P.
PR 22-DEC-1999 99US-01336128P.
PR 23-MAR-1999 99US-01336128P.

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98US-0077791P

98US-0078004P

98US-0078010P

98US-0078910P

98US-0078936P

98US-0078936P

98US-0078938P

98US-0078938P

98US-0079663P

98US-0079664P

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98US-0079664P

98US-008134P

98US-008136P

98US-008137P

98US-0081338P

98US-0081338P

98US-0081338P

98US-008136P

98US-0081338P

98US-0081338P

98US-0081338P

98US-0081338P
 Gaps
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                                                                                                                                                                                                                                                                                                                                               / Match 2.9%; Score 12.4; DB 1; Length 18; Local Similarity 92.9%; Pred. No. 4.3e+02; nes 13; Conservative 0; Mismatches 1; Indels
21-MAR-2000; 2000MO-US007532.
30-MAY-2000; 2000MO-US018439.
17-MAY-2000; 2000MO-US014042.
22-MAY-2000; 2000MO-US014042.
30-MAY-2000; 2000MO-US014941.
30-MAY-2000; 2000MO-US01524.
28-UUL-2000; 2000MO-US02328.
24-AUG-2000; 2000MS-00723749.
37-NOV-2000; 2000MS-00723749.
37-NOV-2000; 2000MS-00723749.
30-DEC-2000; 2000MS-00723749.
30-DEC-2000; 2000MS-00723749.
32-MAR-2001; 2001MS-00816520.
32-MAR-2001; 2001MS-00816620.
32-MAR-2001; 2001MS-00816620.
32-MAR-2001; 2001MS-00816620.
32-MAR-2001; 2001MS-00816744.
32-MAR-2001; 2001MS-00816749.
32-MAY-2001; 2001MS-00816780.
32-MAY-2001; 2001MS-0081790.
31-UNN-2001; 2001MS-00817953.
31-UNN-2001; 2001MS-00816335.
31-UNN-2001; 2001MS-00816335.
31-UNN-2001; 2001MS-0081663.
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97US-0064249P.
97US-0065311P.
98US-0077450P.
98US-0077632P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human PRO DNA PCR primer #6.
                                                                                                                                                                                                                                                                                                                                                                                           215 GAACTCGGTGGCGG 228
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ADB76236 standard; DNA; 18
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03-NOV-1997;
13-NOV-1997;
21-NOV-1997;
11-MAR-1998;
11-MAR-1998;
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Best Local S
Matches 13
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Hillan KJ; Roy MA, Shelton DL;

15-MAY-1998; 18-MAY-1998; 22-MAY-1998; 22-MAY-1998; 22-MAY-1998; 22-MAY-1998; 23-MAY-1998; 24-MAY-1998; 28-MAY-1998; 28-MAY-1998; 28-MAY-1998; 26-JUN-1998; 26-JUN-1998; 20-NUV-1998; 20-NUV-1998; 22-DEC-1998; 23-DEC-1998;

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The present invention relates to the isolation of novel human PRO polypeptides are secreted and transmembrane proteins. The PRO polypeptides are secreted and transmembrane proteins. The PRO polypeptides are useful for detecting other PRO polypeptides, for linking bioactive molecules to cells expressing PRO polypeptides, for modulating biological activities of cells expressing PRO polypeptides, and for toxin, radiolabel or antibody, and cause cell death. The PRO polypeptides are useful for treating neuropathy and neuropathy related diseases such as Charcher-Marie-Tooth disorder, Refenum's diseases, and Krabbe's disease. The polynucleotide sequences encoding PRO polypeptides are useful as hybridisation probes, in chromosome and gene mapping, in the generation of antisense RNA and DNA, in the preparation of PRO polypeptides, for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human, 88; PCR, secreted protein; transmembrane protein; PRO; cytostatic; ophthalmological; antiarthritic; osteopathic; antirheumatic; vulnerary; auditory; tumour growth; retinal disorder; sports-related joint problem; articular cartilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer.
                                                                                 New PRO polypeptides useful for treating peripheral neuropathy, neuropathies associated with systemic disease such as post-polio syndrome or AIDS-associated syndrome.
                                                                                                                                                                                                                                                                                                                                                                                    Query Match

2.9%; Score 12.4; DB 1; Length 18;
Best Local Similarity 92.9%; Pred. No. 4.3e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
PJ, Grimaldi JC, Gurney AL, 1
Napier MA, Pan J, Paoni NF,
Williams PM, Wood WI;
                                                                                                                                                    Example 4; Page 125; 425pp; English
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97US-0065311P.
97US-0065311P.
97US-0077450P.
98US-0077440P.
98US-0077641P.
98US-0077641P.
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98US-0077641P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADC43662 standard; DNA; 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Goddard A, Godowski PJ,
Kljavin IJ, Kuo SS, Na
Stewart TA, Tumas D, W
                                                          WPI; 2003-755118/71.
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10 - MAR-1998;
11 - MAR-1998;
11 - MAR-1998;
12 - WAR-1998;
12 - WAR-1998;
13 - MAR-1998;
20 - MAR-1998;
20 - MAR-1998;
20 - MAR-1998;
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13-NOV-1997
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       Eaton DL;
Gerritsen ME;
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W, Gerber H,
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Gao
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Fong S, (
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2000MO-US003341.
2000MO-US005841.
2000MO-US005841.
2000MO-US005841.
2000MO-US00532.
2000MO-US013705.
2000MO-US013705.
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2000MO-US032678.
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2000MO-US032678.
2000MO-US03678.
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98US-0086622P.
98US-0086414P.
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98US-0086416P.
98US-0087208P.
98US-0087208P.
98US-0091329P.
98US-0091329P.
98US-010038P.
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98US-010038P.
98US-0101329F.
98US-011329F.
98US-011329F.
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99US-0139557P.
99US-013023P.
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02-DEC-1999; 02-DBC-1999; 16-DBC-1999; 30-DBC-1999; 30-DBC-1999; 05-JAN-2000; 06-JAN-2000; 06-JAN-2000; 11-FEB-2000; 18-FEB-2000;

07-JUL-1999; 26-JUL-1999; 28-JUL-1999; 29-OCT-1999; 30-NOV-1999;

05-JAN-1999) 08-MAR-1999) 12-WAR-1999) 12-WAR-1999) 21-APR-1999) 26-APR-1999) 26-APR-1999) 14-MAY-1999) 14-MAY-1999) 16-JUN-1999) 23-JUN-1999)

24-FEB-2000; 02-MAR-2000; 21-MAR-2000; 21-MAR-2000; 30-MAR-2000; 17-MAY-2000; 30-MAY-2000; 30-MAY-2000; 28-JUL-2000; 24-AUG-2000;

22-MAR-2001; 2 25-MAY-2001; 2 01-JUN-2001; 2 20-JUN-2001; 2

Ashkenazi AJ, Ferrara N, F

30-JUL-2001;

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Gaps ö

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PR 20-WAR-1998; 98US-007939P.
PR 25-WAR-1998; 98US-0079564P.
PR 27-WAR-1998; 98US-0079564P.
PR 27-WAR-1998; 98US-0079664P.
PR 27-WAR-1998; 98US-0079664P.
PR 31-WAR-1998; 98US-0079786P.
PR 31-WAR-1998; 98US-0079786P.
PR 31-WAR-1998; 98US-0079786P.
PR 31-WAR-1998; 98US-0079320P.
PR 31-WAR-1998; 98US-0080107P.
PR 31-WAR-1998; 98US-0080107P.
PR 31-WAR-1998; 98US-0080137P.
PR 01-APR-1998; 98US-0080137P.
PR 01-APR-1998; 98US-0080137P.
PR 01-APR-1998; 98US-0080137P.
PR 01-APR-1998; 98US-008132P.
PR 15-APR-1998; 98US-008132P.
PR 22-APR-1998; 98US-00813P.
PR 22-APR-1998; 98US-008132P.
PR 22-APR-1998; 98US-008133P.
PR 22-APR-1998; 98US-00
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PR 22-MAY-1998; 98US-0086486 P. PR 28-MXY-1998; 98US-008708 P. PR 28-MXY-1998; 98US-008708 P. PR 28-MXY-1998; 98US-008708 P. PR 28-MXY-1998; 98US-00106413 P. PR 26-UJN-1998; 98US-00106413 P. PR 26-UJN-1998; 98US-00106413 P. PR 26-UJN-1998; 98US-001064141 P. PR 26-UJN-1998; 98US-001064141 P. PR 26-UJN-1998; 98US-00106461 P. PR 26-UJN-1998; 98US-0010637 P. PR 26-UJN-1998; 98US-0010637 P. PR 20-DEC-1998; 98US-0010631 P. PR 20-DEC-1998; 98US-0010637 P. PR 20-DEC-1998; 98US-0010631 P. PR 20-DEC-1998; 98US-0010637 P. PR 20-DEC-1998; 98US-0010631 P. PR 20-DEC-1999; 98US-0010637 P. PR 20-DEC-1999; 98US-0010631 P. PR 20-DEC-1999; 98US-0010637 P. PR 20-DEC-1999; 98US-0010631 P. PR 20-DEC-2000; 2000MO-US001231 P. PR 20-DEC-

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ADC61422 Standard; DNA; 18 BP.
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ADC61422;
XX
T 18-DEC-2003 (first entry)
XX
Human; 8s; PCR; secreted protein
XW
Wyhthalmological; antiarthritic;
XX
Human; 8s; PCR; secreted protein
XW
Wyhthalmological; antiarthritic;
XX
Auditory; tumour growth; retinal
XX
XX
Homo sapiens.
XX
XX
YS
Homo sapiens.
XX
XX
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Homo sapiens.
XX
XX
YS
13-MAR-2003,
XX
XX
XI
10-MAR-1997; 97US-0064249P.
PR 13-MAR-1997; 97US-006434P.
PR 13-MAR-1998; 98US-0077450P.
PR 11-MAR-1998; 98US-0077641P.
PR 11-MAR-1998; 98US-0077641P.
PR 11-MAR-1998; 98US-0077691P.
PR 12-MAR-1998; 98US-0077691P.
PR 12-MAR-1998; 98US-0077691P.
PR 12-MAR-1998; 98US-0077691P.
PR 20-MAR-1998; 98US-0077665P.
PR 27-MAR-1998; 98US-0077669P.
PR 27-MAR-1998; 98US-007769P.
PR 27-MAR-1998; 98US-0077
22-MAR-2001, 2001US-00816744.
22-MAR-2001, 2001US-00816920.
22-MAR-2001, 2001US-00854208.
10-MAY-2001, 2001US-00854208.
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01-JUN-2001, 2001WO-US017092.
01-JUN-2001, 2001WO-US017800.
05-JUN-2001, 2001US-00874503.
14-JUN-2001, 2001US-00882636.
19-JUN-2001, 2001WS-0US019692.
20-JUN-2001, 2001WO-US019692.
20-JUN-2001, 2001WO-US01966.
09-JUL-2001, 2001WO-US01966.
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18 GAACTCCGTGGCGG 5
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12-APR-1999; 99US-00130232P.
21-APR-1999; 99US-0131022P.
28-APR-1999; 99US-01314287P.
14-MAY-1999; 99US-01314287P.
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14-MAY-1999; 99US-01314287P.
14-MAY-1999; 99US-01314287P.
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15-JUN-1999; 99US-0131352-7P.
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25-AUG-1999; 99US-0141037P.
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25-AUG-1999; 99US-014623855.
26-ANA-2000; 2000WO-US00376.
21-MAR-2000; 2000WO-US0036319.
21-MAR-2000; 2000WO-US01341.
22-MAY-2000; 2000WO-US01341.
23-MAY-2000; 2000WO-US01341.
24-AUG-2000; 2000WO-US01341.
25-MAY-2000; 2000WO-US01341.
26-MAY-2000; 2000WO-US01341.
27-MAY-2000; 2000WO-US01341.
28-MAY-2000; 2000WO-US01341.
28-MAY-2000; 2000WO-US01341.
28-MAY-2000; 2000WO-US01341.
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20-MAY-2000; 2000WO-US01341.
20-MAY-2000; 2000WO-US01341.
20-MAY-2000; 2000WO-US01341.
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99US-0108621P.
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112-MAR-1999,
12-MAR-1999,
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27-MAR-1998; 27-MAR-1998; 27-MAR-1998;

27-MAR-1998; 27-MAR-1998; 30-MAR-1998; 30-MAR-1998; 31-MAR-1998; 31-MAR-1998; 31-MAR-1998;

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Human, ss; PCR; secreted protein; transmembrane protein; PRO; cytostatic; ophthalmological; antiarthritic; osteopathic; antirheumatic; vulnerary; auditory; tumour growth, retinal disorder; sports-related joint problem; articular cartilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer.
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2.9%; Score 12.4; DB 1;
Best Local Similarity 92.9%; Pred. No. 4.3e+02;
Matches 13; Conservative 0; Mismatches 1;
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97US-0064349P-
97US-0066364P-
98US-0077451P-
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98US-0078910P-
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98US-0079788P-
05-JUN-2001; 2001US-00874503.
14-JUN-2001; 2001US-00882636.
19-JUN-2001; 2001US-0086542.
20-JUN-2001; 2001WO-US021066.
29-JUN-2001; 2001WO-US021066.
09-JUL-2001; 2001US-00918585.
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03-NOV-1997;
21-NOV-1997;
21-NOV-1999;
11-MAR-1998;
11-MAR-1998;
12-MAR-1998;
12-MAR-1998;
13-MAR-1998;
17-MAR-1998;
20-MAR-1998;
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802-00803378805-00803338805-00803338805-00810498805-00810708805-00810708805-00811958	605 - 00812037 605 - 00812037 605 - 00818129 805 - 0081819 805 - 00819528 805 - 00819528 805 - 00819528 805 - 00826698 805 - 00827008 805 - 00827008 805 - 00827008 805 - 00827008	803 - 0083336 P	805-00846098 805-00846098 805-00846279 805-00846379 805-00846378 805-00853389 805-00853389 805-00853389 805-008553889 805-00855898	980S - 0086392P 980S - 0086392P 980S - 0086392P 980S - 0086434P 980S - 0086486F 980S - 0086486F 980S - 0087098P 980S - 0087208P 980S - 0097098P 980S - 0097010P 980S - 0097010P
1-APR-1998 1-APR-1998 1-APR-1998 8-APR-1998 8-APR-1998 8-APR-1998	9-APR-1998 9-APR-1998 5-APR-1998 5-APR-1998 1-APR-1998 1-APR-1998 1-APR-1998 1-APR-1998 2-APR-1998	7. APR-1998 B. APR-1998 B. APR-1998 P. APR-1998 P. APR-1999 P. APR-1999 P. APR-1999 P. APR-1999 P. APR-1999 P. APR-1998 P. APR-1998 P. APR-1998	7. WAY-1998 7. WAY-1998 7. WAY-1998 7. WAY-1998 8. WAY-1998	15-7647.1998; 22-MAY-1998; 22-MAY-1998; 22-MAY-1998; 22-MAY-1998; 22-MAY-1998; 28-MAY-1998; 26-UN-1998; 26-UN-1998
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PR 20-NOV-1996 98US-0109304P
PR 20-NOV-1996 98US-0109304P
PR 22-DEC-1996 98US-010202656
PR 22-DEC-1996 98US-010202656
PR 22-DEC-1996 98US-010202656
PR 22-DEC-1996 98US-010202656
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PR 25-MAR-2000 2000MO-US001313
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PR 25-MAR-2000 2000MO-US001313
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PR 27-MAR-2000 2000MO-US001313
PR 27-MAR-2000 2000MO-US0013232
PR 27-MAR-2000 2000MO-US013232
PR 27-MAR-2001 2000MO-

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                                                                       Gaps
                                                                        ;
                                                Score 12.4; DB 1; Length 18;
Pred. No. 4.3e+02;
0; Mismatches 1; Indels
                                                                      1; Indels
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 30-JUL-2001; 2001US-00918585.
                                               2.9%;
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                                                                                         215 GAACTCGGTGGCGG 228
                                                                                                                                                                                                                          Human PRO 274 PCR primer #4
                                                                                                                                                               ADC66486 standard; DNA; 18
                                                                                                                                                                                                       18-DEC-2003 (first entry)
                                                                                                             18 daacrccgreeces 5
                                                           Best Local Similarity 92.9
Matches 13; Conservative
                    (GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
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20-MAR-1998;
20-MAR-1998;
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                                                                                                                                                                                   ADC66486;
                                                   Query Match
                                                                                                                                            RESULT 697
                                                                                                                                                      ADC66486,
# X # X
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Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Baton DL; Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME; Goddard A, Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ; Kljavin IJ, Kwo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton DL; Stewart TA, Tumas D, Williams PM, Wood WI;
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28-JUL-2000; 2000MO-US012564.
24-AUC-2000; 2000MO-US02338.
8-NOV-2000; 2000MS-00723749.
27-NOV-2000; 2000MS-00723749.
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20-DEC-2000; 2000MS-00747259.
22-MAR-2001; 2001MS-00816744.
22-MAR-2001; 2001MS-00816520.
22-MAR-2001; 2001MS-00816744.
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9903-00380142.
30-DEC-1999)
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02-DEC-1999)
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10-MAY-2001; 2001US-00854280.

25-MAY-2001; 2001US-00854280.

01-UNA-2001; 2001US-0085038.

01-UNA-2001; 2001US-0087600.

05-UNA-2001; 2001US-0087600.

14-UNA-2001; 2001US-00886342.
99WO-US000106.
99US-00254465.
99WO-US00528.
99US-00265686.
99WO-US005190.
                                                                                                                                     99US-00284291.
99US-00311832.
99WO-US010733.
99WO-US012252.
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29-JUN-2001; 2001WO-US021066
09-JUL-2001; 2001WO-US021735
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30-MAY-2000, 2000WO-US014941
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05-7AN-1999;
05-MAR-1999;
06-MAR-1999;
10-MAR-1999;
110-MAR-1999;
112-MAR-1999;
14-MAY-1999;
14-MAY-1999;
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Novel secreted and transmembrane polypeptides and polynucleotides encoding them, useful for treating wound healing, tissue growth and muscle generation and regeneration, amyotrophic lateral sclerosis or neuropathy.

Example 4; SEQ ID NO 14; 472pp; English.